

GenCore version 5.1.3
Copyright (c) 1993 - 2002 CompuGen Ltd.

OM nucleic - nucleic search, using sw model

Run on: December 24, 2002, 16:23:56 ; Search time 206.404 Seconds
(without alignments)
13583.723 Million cell updates/sec

Title: US-09-708-724A-1

Perfect score: 1245

Sequence: 1 atgggaccctgctcagtggt.....ccaggaagcgggctgagtag 1245

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 2185239 seqs, 1125999159 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

N_Geneseq_101002.*
1: /SID22/gcgdata/geneseq/geneseq-emb1/NA1980.DAT.*
2: /SID22/gcgdata/geneseq/geneseq-emb1/NA1981.DAT.*
3: /SID22/gcgdata/geneseq/geneseq-emb1/NA1982.DAT.*
4: /SID22/gcgdata/geneseq/geneseq-emb1/NA1983.DAT.*
5: /SID22/gcgdata/geneseq/geneseq-emb1/NA1984.DAT.*
6: /SID22/gcgdata/geneseq/geneseq-emb1/NA1985.DAT.*
7: /SID22/gcgdata/geneseq/geneseq-emb1/NA1986.DAT.*
8: /SID22/gcgdata/geneseq/geneseq-emb1/NA1987.DAT.*
9: /SID22/gcgdata/geneseq/geneseq-emb1/NA1988.DAT.*
10: /SID22/gcgdata/geneseq/geneseq-emb1/NA1989.DAT.*
11: /SID22/gcgdata/geneseq/geneseq-emb1/NA1990.DAT.*
12: /SID22/gcgdata/geneseq/geneseq-emb1/NA1991.DAT.*
13: /SID22/gcgdata/geneseq/geneseq-emb1/NA1992.DAT.*
14: /SID22/gcgdata/geneseq/geneseq-emb1/NA1993.DAT.*
15: /SID22/gcgdata/geneseq/geneseq-emb1/NA1994.DAT.*
16: /SID22/gcgdata/geneseq/geneseq-emb1/NA1995.DAT.*
17: /SID22/gcgdata/geneseq/geneseq-emb1/NA1996.DAT.*
18: /SID22/gcgdata/geneseq/geneseq-emb1/NA1997.DAT.*
19: /SID22/gcgdata/geneseq/geneseq-emb1/NA1998.DAT.*
20: /SID22/gcgdata/geneseq/geneseq-emb1/NA1999.DAT.*
21: /SID22/gcgdata/geneseq/geneseq-emb1/NA2000.DAT.*
22: /SID22/gcgdata/geneseq/geneseq-emb1/NA2001A.DAT.*
23: /SID22/gcgdata/geneseq/geneseq-emb1/NA2001B.DAT.*
24: /SID22/gcgdata/geneseq/geneseq-emb1/NA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
C 1	162.2	13.0	1872	24	ABL99794 Human secretory po
2	160	12.9	143068	21	AAF21105 Human low adenosin
3	160	12.9	143068	21	AAF21272 Human low adenosin
4	160	12.9	143068	21	AAA34983 Human adenosine re
5	160	12.9	143068	21	AAA35150 Human adenosine re
6	160	12.9	143068	24	AA658124 Ovary cancer relat
7	160	12.9	149412	21	AAA35151 Human adenosine re
8	160	12.9	152740	21	AAF21273 Human low adenosin
9	159	12.8	369	23	AA568798 DNA encoding novel

10	159	12.8	401	22	ABA72549	Human foetal liver
11	159	12.8	401	22	AAK20973	Human brain expres
12	159	12.8	401	22	AAK47126	Human bone marrow
13	159	12.8	401	22	AAI52962	Probe #21648 used
14	159	12.8	401	24	ABS21353	Human genome-deriv
15	159	12.8	552	22	ABA60008	Human foetal liver
16	159	12.8	552	22	AAK08279	Human brain expres
17	159	12.8	552	22	AAK34158	Human bone marrow
18	159	12.8	552	22	AAI39881	Probe #8567 used t
19	159	12.8	552	24	ABS08919	Human genome-deriv
20	159	12.8	1002	23	AA588799	DNA encoding novel
21	159	12.8	1683	23	AA585679	DNA encoding novel
22	154.2	12.4	4977	22	AA526628	Human genomic DNA
C 23	102.4	8.2	526	22	AAI35808	Human musculoskele
C 24	97.8	7.9	2212	23	AA578334	DNA encoding novel
C 25	89.8	7.2	527	22	AAI35807	Human musculoskele
C 26	89.8	7.2	527	22	AAI35807	Human musculoskele
C 27	87	7.0	1019	22	AAH98277	Human EST-derived
28	85.8	6.9	2750	21	AAC69110	Human secreted pro
29	85.8	6.9	2752	21	AAC69119	Human secreted pro
30	60	4.8	759	22	AAF68303	Human lung tumour
31	60	4.8	759	24	ABK38214	CDNA encoding clon
32	52	4.2	893	23	AA565934	DNA encoding novel
33	52	4.2	1503	23	AA578107	DNA encoding novel
34	51.2	4.1	514	22	ABA60429	Human foetal liver
35	51.2	4.1	514	22	AAK08707	Human brain expres
36	51.2	4.1	514	22	AAK34594	Human bone marrow
37	51.2	4.1	514	22	AAI40313	Probe #8999 used t
C 38	44	3.5	77536	21	AAAI4651	Nucleotide sequenc
39	42.8	3.4	1026	23	AA578332	DNA encoding novel
C 40	42.6	3.4	707	24	ABQ39154	Oligonucleotide fo
41	42.6	3.4	707	24	ABQ39155	Oligonucleotide fo
42	41.2	3.3	461	23	ABV33700	Human prostate exp
43	41.2	3.3	461	23	ABV42603	Human prostate exp
44	40.6	3.3	438	23	ABV03397	Human prostate exp
45	40.6	3.3	1278	20	AA590824	DNA encoding human

ALIGNMENTS

RESULT 1

ABL99794/C

ID ABL99794 standard; cDNA: 1872 BP.

XX ABL99794;

AC ABL99794;

XX 03-OCT-2002 (first entry)

DT Human secretory polynucleotide (sptm) 49.

DE Human; ss; gene; secretory protein; secretory polynucleotides; SPTM;

XX SPTM-related disease; somatic gene therapy; germline gene therapy;

KW severe combined immunodeficiency; intracellular parasite protection;

KW fungal parasite; protozoan parasite; cell proliferative disorder; cancer;

KW immune disorder; AIDS; neurological disorder; Parkinson's disease;

KW motor neuron disorder; demyelinating disease; multiple sclerosis;

KW meningitis; abscess; prion diseases; cerebral palsy;

KW neuroskeletal disorder; peripheral nervous system disorder;

KW dermatomyositis; polymyositis; myopathy; myasthenia gravis;

KW mental disorder; Tourette's syndrome.

XX Homo sapiens.

OS WO200220756-A2.

XX 14-MAR-2002.

PD 30-AUG-2001; 2001WO-US27297.

XX 05-SEP-2000; 2000US-229747P.

XX 05-SEP-2000; 2000US-229748P.

PR 05-SEP-2000; 2000US-229749P.

CC and/or bronchoconstriction) and/or lung inflammation, allergy(ies)
 CC and/or surfactant hypoproduction which are associated with a disease or
 CC condition selected from pulmonary vasoconstriction, inflammation,
 CC allergies, asthma, impaired respiration, respiratory distress syndrome
 CC (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary
 CC hypertension, emphysema, chronic obstructive pulmonary disease (COPD),
 CC pulmonary transplantation rejection, pulmonary infections, bronchitis,
 CC and/or cancer. AAF18434 to AAF21543 represent human polynucleotide
 CC fragments and antisense oligonucleotides used in the exemplification of
 CC the present invention.

XX
 SQ Sequence 143068 BP; 41194 A; 30122 C; 32403 G; 39349 T; 0 other;
 Query Match 12.9%; Score 160; DB 21; Length 143068;
 Best Local Similarity 82.1%; Pred. No. 1.7e-33;
 Matches 184; Conservative 0; Mismatches 40; Indels 0; Gaps 0;
 QY 707 CTGAATCTCAAAATTTTGAAGAATCTTTTGGTCCACACACACCAAGAAATAATA 766
 DB 21653 CTGAATCTCAAAATTTTGAAGAATTTGTCTACTACAGCTCCAATTTGAATATAA 21712
 QY 767 AACAGGAGGAGGAGGATGAATTTGGCTTACACACCCCTCCAGTACGAGAAACACCTG 826
 DB 21713 AACAGGAGGAGGAGGATGAATTTGGCTTACACCCCTCCAGTACGAGAAACATCTG 21772
 QY 827 TACCATCTCTTCAGTACAGAAATAGAGACCCCTCCAGTACGAGAAACATCTG 886
 DB 21773 TACCATCTCTTCAGTACGAGAAATAGAGATCCCAAGAAATTTATGCTCTGCTG 21832
 QY 887 CCATAGCTGGAGAGCCCTTAGGACATTCGACTTTTCACTATTCT 930
 DB 21833 TCATAGCTGGAGAGCCCTTAGGACCTTGTGCTTTTCTCTATTCT 21876

RESULT 3
 AAF21272
 ID AAF21272 standard; DNA; 143068 BP.
 AC AAF21272;
 XX
 XX 14-MAR-2001 (first entry)
 XX Human low adenosine antisense oligonucleotide related sequence #2839.
 DE
 XX

Low adenosine antisense oligonucleotide; phosphorothioate; allergy;
 KW human; airway disorder; bronchoconstriction; lung inflammation;
 KW surfactant depletion; respiratory; bronchodilator; antiinflammatory;
 KW immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic;
 KW respiratory obstruction; pulmonary obstruction; impeded respiration;
 KW surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS;
 KW respiratory distress syndrome; pulmonary fibrosis; allergic rhinitis;
 KW pulmonary hypertension; emphysema; pain; cystic fibrosis; impaired airway;
 KW chronic obstructive pulmonary disease; pulmonary transplantation rejection;
 KW cancer; ss.

XX Homo sapiens.
 OS
 XX WO200062736-A2.
 PN
 XX 26-OCT-2000.
 PD
 XX 24-MAR-2000; 2000WO-US08020.
 XX
 PF
 XX 06-APR-1999; 99US-0127958.
 PR
 XX (UYEC-) UNIV EAST CAROLINA.
 PA (NYCE/) NYCE J W.
 XX
 XX NYCE JW;
 PI
 XX WPI; 2000-679539/66.
 DR
 XX Low adenosine (A) content antisense oligonucleotides which do not

PT trigger adenosine receptors during metabolism, useful e.g. for treating
 PT cancers and respiratory obstructions -
 XX
 PS Disclosure; Page 1186-1219; 1592pp; English.

XX The present invention describes low adenine (A) content antisense
 CC oligonucleotides and compositions (I) comprising them. In the antisense
 CC oligonucleotides the A is replaced by a 'Universal' or alternative base.
 CC (I) can have respiratory, bronchodilator, antiinflammatory, analgesic,
 CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities.
 CC The antisense oligonucleotides and (I) can be used to down-regulate the
 CC expression and/or activity of target polypeptides associated with
 CC lung/respiratory disorders and malignancies, such as stimulating and
 CC activating peptide factors and transmitters, transcription factors,
 CC immunoglobulins and antibodies, antibody receptors, cytokines and
 CC chemokines, endogenously produced specific and non-specific enzymes,
 CC binding proteins, adhesion molecules and their receptors, cytokine and
 CC chemokine receptors, adenosine receptors, bradykinin receptors, central
 CC nervous system (CNS) and peripheral nervous and non-nervous system
 CC receptors, CNS and peripheral nervous and non-nervous system peptide
 CC transmitters, defensins, growth factors, vasoactive peptides and
 CC receptors, binding proteins and malignancy associated proteins. The
 CC antisense oligonucleotides may be used in this way to treat disorders
 CC including respiratory obstruction (especially pulmonary obstruction
 CC and/or bronchoconstriction) and/or lung inflammation, allergy(ies)
 CC and/or surfactant hypoproduction which are associated with a disease or
 CC condition selected from pulmonary vasoconstriction, inflammation,
 CC allergies, asthma, impeded respiration, respiratory distress syndrome
 CC (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary
 CC hypertension, emphysema, chronic obstructive pulmonary disease (COPD),
 CC pulmonary transplantation rejection, pulmonary infections, bronchitis,
 CC and/or cancer. AAF18434 to AAF21543 represent human polynucleotide
 CC fragments and antisense oligonucleotides used in the exemplification of
 CC the present invention.

XX
 SQ Sequence 143068 BP; 41194 A; 30122 C; 32403 G; 39349 T; 0 other;
 Query Match 12.9%; Score 160; DB 21; Length 143068;
 Best Local Similarity 82.1%; Pred. No. 1.7e-33;
 Matches 184; Conservative 0; Mismatches 40; Indels 0; Gaps 0;
 QY 707 CTGAATCTCAAAATTTTGAAGAATCTTTTGGTCCACACACCAAGAAATAATA 766
 DB 21653 CTGAATCTCAAAATTTTGAAGAATTTGTCTACTACAGCTCCAATTTGAATATAA 21712
 QY 767 AACAGGAGGAGGAGGATGAATTTGGCTTACACACCCCTCCAGTACGAGAAACACCTG 826
 DB 21713 AACAGGAGGAGGAGGATGAATTTGGCTTACACCCCTCCAGTACGAGAAACATCTG 21772
 QY 827 TACCATCTCTTCAGTACAGAAATAGAGACCCCTCCAGTACGAGAAATTCGCGACTGCTA 886
 DB 21773 TACCATCTCTTCAGTACGAGAAATAGAGATCCCAAGAAATTTATGCTCTGCTG 21832
 QY 887 CCATAGCTGGAGAGCCCTTAGGACATTCGACTTTTCACTATTCT 930
 DB 21833 TCATAGCTGGAGAGCCCTTAGGACCTTGTGCTTTTCTCTATTCT 21876

RESULT 4
 AAA34983
 ID AAA34983 standard; DNA; 143068 BP.
 AC AAA34983;
 XX
 XX 28-JUL-2000 (first entry)
 DT Human adenosine receptor related polynucleotide SEQ ID NO:2672.
 DE
 XX
 XX Human; adenosine receptor; low adenine antisense oligonucleotide;
 KW phosphorothioate; impaired respiration; inflammation; allergy;
 KW allergic disease; bronchoconstriction; inhibitor; antiinflammatory;
 KW antiasthmatic; antiasthmatic; cytostatic; analgesic; impaired airway;
 KW lung disease; ischaemic condition; pulmonary vasoconstriction; asthma;

XX 28-JUL-2000 (first entry)
XX Human adenosine receptor related polynucleotide 2nd SEQ ID NO:25.
XX
XX
XX Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytotatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
XX
XX Homo sapiens.
XX WO200009525-A2.
XX
XX 24-FEB-2000.
XX
XX 03-AUG-1999; 99WO-US17712.
XX
XX 03-AUG-1998; 98US-0095212.
XX
XX (UYEC-) UNIV EAST CAROLINA.
XX
XX Nyce JW;
XX
XX WPI; 2000-205971/18.
XX
XX New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or cancers -
XX
XX Disclosure; Page 1138-1171; 1343pp; English.
XX
XX The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytotatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, impaired respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasize to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32313 to AAA35312 represent the nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA32323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences given in the sequence listing.
XX
XX Sequence 149412 BP; 43049 A; 31388 C; 33852 G; 41123 T; 0 other;
XX
XX Query Match 12.9%; Score 160; DB 21; Length 149412;
XX Best Local Similarity 82.1%; Pred. No. 1.8e-33;
XX Matches 184; Conservative 0; Mismatches 40; Indels 0; Gaps 0;
XX
XX 707 CTGAATCTCAAAATTTGGAAGATCTTTGTCCACCCACACCAAGAAATAATA 766
XX
XX 27997 CTGAATCTCAAAATTTGGAAGATTTGTCTTACTACAGCTCCAAATGATATAAAA 28056

QY 767 AACAGAGAGGAGGATGAAATTTGGGCTCTACACCCCTCCAGTAGCAGAAACACCTG 826
DB 28057 AACAGAGAGGAGGATGAAATTTGGGCTCTATACCGCTCTCCAGATGCAAGACATCTG 28116
QY 827 TACCATCTCTTCAGTAAACAGAAATAGAGACCCCACTGCAAGAATTCGCGGACTGCTA 886
DB 28117 TACCATCTCTTCAGTGGCAGAAATAGAGATCCAGTACAAAGAATTTTATGCTCTGCTG 28176
QY 887 CCATAGCTGGAGAGCCCTTAGGACATTTGCACATTTTCTTCTTCTTCTTCTTCTTCTTCT 930
DB 28177 TCATAGCTGGAGAGCCCTTAGGACCTTGTGCTTTTCTTCTTCTTCTTCTTCTTCTTCT 28220
RESULT 8
AAF21273
ID AAF21273 standard; DNA; 152740 BP.
XX
AC AAF21273;
XX
DT 14-MAR-2001 (first entry)
XX
DE Human low adenosine antisense oligonucleotide related sequence #2840.
XX
KW Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytotatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis; cancer; ss.
XX
OS Homo sapiens.
XX
PN WO200062736-A2.
XX
PD 26-OCT-2000.
XX
PF 24-MAR-2000; 2000WO-US08020.
XX
PR 06-APR-1999; 99US-0127958.
XX
PA (UYEC-) UNIV EAST CAROLINA.
XX
XX (NYCE/) NYCE J W.
XX
XX Nyce JW;
XX
XX WPI; 2000-679539/66.
XX
PT Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions -
XX
XX Disclosure; Page 1219-1254; 1592pp; English.
XX
XX The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytotatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and chemokine receptors, adenosine receptors, bradykinin receptors, central nervous system (CNS) and peripheral nervous and non-nervous system receptors, CNS and peripheral nervous and non-nervous system peptide transmitters, defensins, growth factors, vasoactive peptides and

CC receptors, binding proteins and malignancy associated proteins. The
 CC antisense oligonucleotides may be used in this way to treat disorders
 CC including respiratory obstruction (especially pulmonary obstruction
 CC and/or bronchoconstriction) and/or lung inflammation, allergy(ies)
 CC and/or surfactant hypoproduction which are associated with a disease or
 CC condition selected from pulmonary vasoconstriction, inflammation,
 CC allergies, asthma, impeded respiration, respiratory distress syndrome
 CC (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary
 CC hypertension, emphysema, chronic obstructive pulmonary disease (COPD),
 CC pulmonary transplantation rejection, pulmonary infections, bronchitis,
 CC and/or cancer. AAF18434 to AAF21543 represent human polynucleotide
 CC fragments and antisense oligonucleotides used in the exemplification of
 CC the present invention.

XX SQ Sequence 152740 BP; 44169 A; 32023 C; 34549 G; 41999 T; 0 other;
 Query Match 12.9%; Score 160; DB 21; Length 152740;
 Best Local Similarity 82.1%; Pred. No. 1.8e-33;
 Matches 184; Conservative 0; Mismatches 40; Indels 0; Gaps 0;

Qy 707 CTGAATCTCAAAATTTTGAAGAATCTTTTGTCCCAACACACCCCAAGAAAATAATA 766
 Db 27997 CTGATCTCAAAATTTGGAAGAATTTGTCTCTACTACAGCTCAATTTGAATATAAA 28056

Qy 767 AACAGGAGGGAGGATGAAATTTGGCGTCTACACACCCCTCCAGTAGCAGAAACACCTG 826
 Db 28057 AACAGGAGGGAGGATGAAATTTGGCGTCTATACCGCTCTCCAGATGCAGAAACATCTG 28116

Qy 827 TACCATCTCTTCAGTACAGAAATAGAGACCCCACTGCAAGAAATTCGGCGGACTGCTA 886
 Db 28117 TACCATCTCTCTTCAGTACAGAAATAGAGATCCCACTGCAAGAAATTTATGCTCTGCTG 28176

Qy 887 CCATAGCTGGAGAGCCCTTAGGACATTCGACATTTTCACATTTTCT 930
 Db 28177 TCATAGCTGGAGAGCCCTTAGGACATTTGCTTTTCCCTATTCT 28220

RESULT 9
 AAS68798
 ID AAS68798 standard; cDNA; 369 BP.

XX AC AAS68798;
 XX DT 13-FEB-2002 (first entry)
 XX DE DNA encoding novel human diagnostic protein #4602.

XX Human; chromosome mapping; gene mapping; gene therapy; forensic;
 KW food supplement; medical imaging; diagnostic; genetic disorder; ss.
 XX OS Homo sapiens.

XX PN WO200175067-A2.
 XX PD 11-OCT-2001.

XX PF 30-MAR-2001; 2001WO-US08631.
 XX PR 31-MAR-2000; 2000US-0540217.
 XX PR 23-AUG-2000; 2000US-0649167.

XX PA (HYSE-) HYSEQ INC.
 XX PI Drmanac RT, Liu C, Tang YT;
 XX DR WPI; 2001-639362/73.
 XX DR P-PSDB; ABG04611.

XX PT New isolated polynucleotide and encoded polypeptides, useful in
 PT diagnostics, forensics, gene mapping, identification of mutations
 PT responsible for genetic disorders or other traits and to assess
 PT biodiversity -
 XX

PS Claim 1; SEQ ID No 4602; 103pp; English.

XX The invention relates to isolated polynucleotide (I) and
 CC polypeptide (II) sequences. (I) is useful as hybridisation probes,
 CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome
 CC and gene mapping, and in recombinant production of (II). The
 CC polynucleotides are also used in diagnostics as expressed sequence tags
 CC for identifying expressed genes. (I) is useful in gene therapy techniques
 CC to restore normal activity of (II) or to treat disease states involving
 CC (II). (II) is useful for generating antibodies against it, detecting or
 CC quantitating a polypeptide in tissue, as molecular weight markers and as
 CC a food supplement. (II) and its binding partners are useful in medical
 CC imaging of sites expressing (II). (I) and (II) are useful for treating
 CC disorders involving aberrant protein expression or biological activity.
 CC The polypeptide and polynucleotide sequences have applications in
 CC diagnostics, forensics, gene mapping, identification of mutations
 CC responsible for genetic disorders or other traits to assess biodiversity
 CC and to produce other types of data and products dependent on DNA and
 CC amino acid sequences. AAS64197-AAS94564 represent novel human
 CC diagnostic coding sequences of the invention.

CC Note: The sequence data for this patent did not appear in the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pct_sequences.

XX SQ Sequence 369 BP; 112 A; 92 C; 64 G; 101 T; 0 other;
 Query Match 12.8%; Score 159; DB 23; Length 369;
 Best Local Similarity 82.1%; Pred. No. 1.6e-34;
 Matches 183; Conservative 0; Mismatches 40; Indels 0; Gaps 0;

Qy 707 CTGAATCTCAAAATTTTGAAGAATCTTTTGTCCCAACACACCCCAAGAAAATAATA 766
 Db 99 CTGAGTCTAAATTTCTGAAGAATCTGTGTCCCAACACACCCCAAGAAAATAATA 158

Qy 767 AACAGGAGGGAGGATGAAATTTGGCGTCTACACACCCCTCCAGTAGCAGAAACACCTG 826
 Db 159 AACAGGAGGGAGGATGAAATTTGGCGTCTATACCGCTCTCCCAATCGCAGAAACATCTG 218

Qy 827 TACCATCTCTTCAGTACAGAAATAGAGACCCCACTGCAAGAAATTCGGCGGACTGCTA 886
 Db 219 TACTGCTCTCTTCAGTACAGAAATAGAGACCCCACTGCAAGAAATTTATGCTCTGCTG 278

Qy 887 CCATAGCTGGAGAGCCCTTAGGACATTCGACATTTTCACATTTTCT 929
 Db 279 CCATAGCTGGAGAGCCCTTAGGACATTCGACATTTTCTTCTATTTC 321

RESULT 10
 ABA72549
 ID ABA72549 standard; DNA; 401 BP.

XX AC ABA72549;
 XX DT 01-FEB-2002 (first entry)
 XX DE Human foetal liver single exon nucleic acid probe #20854.

XX KW Human; foetal liver; gene expression; single exon nucleic acid probe; ss.
 XX OS Homo sapiens.

XX PN WO200157277-A2.
 XX PD 09-AUG-2001.

XX PF 30-JAN-2001; 2001WO-US00669.
 XX PR 04-FEB-2000; 2000US-0180312.
 XX PR 26-MAY-2000; 2000US-0207456.
 XX PR 30-JUN-2000; 2000US-0608408.
 XX PR 03-AUG-2000; 2000US-0632366.
 XX PR 21-SEP-2000; 2000US-0234687.
 XX PR 27-SEP-2000; 2000US-0236359.

```
PR 04-OCT-2000; 2000GB-0024263.
PA (MOLE-) MOLECULAR DYNAMICS INC.
PI Penn SG, Hanzel DK, Chen W, Rank DR;
XX WPI; 2001-483447/52.
XX Human genome-derived single exon nucleic acid probes useful for
PT analyzing gene expression in human fetal liver -
XX
PS Claim 4; SEQ ID NO 20854; 639pp + sequence listing; English.
CC The invention relates to a single exon nucleic acid probe for
CC measuring human gene expression in a sample derived from human foetal
CC liver. The single exon nucleic acid probes may be used for predicting,
CC measuring and displaying gene expression in samples derived from human
CC fetal liver. The present sequence is a single exon nucleic acid
CC probe of the invention.
CC Note: The sequence data for this patent did not form part of the
CC printed specification, but was obtained in electronic format directly
CC from WIPO at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 401 BP; 115 A; 92 C; 87 G; 107 T; 0 other;
Query Match 12.8%; Score 159; DB 22; Length 401;
Best Local Similarity 82.1%; Pred. No. 1.7e-34;
Matches 183; Conservative 0; Mismatches 40; Indels 0; Gaps 0;
QY 707 CTGATCTCAAAATTTGAAGAATCTTTTGTCCACCCACACACCCCAAGAAATAATA 766
DB 161 CTGAGCTCTAAATCTGAAGAATCTGTGTCCACCCACACAGCTTCAATTGAAATAAAA 220
QY 767 AACAGGAGGGGAGGATGAAATTTGGCGTCTACCAACCCCTCCAGTAGCAGAAACACCTG 826
DB 221 AACAGGAGGGGAGGATGAAATTTGGCGTCTACCAACCCCTCCAGTAGCAGAAACATCTG 280
QY 827 TACCACTCTCTTCAGTAACAGAAATAGAGACCCCACTGCAAAAGAAATTCGCGGACTGCTA 886
DB 281 TGCCGCTCTCTCGGTAGCAGGAATAGAGACCCCAATACAAAGAAATTTACGCTCTGCTG 340
QY 887 CCATAGCTGGAGAGCCCTTAGGACATTTGCACATTTTCACTATTTC 929
DB 341 CCATAGCTGGAGAGCCCTTAGGACATTTGCACATTTTCACTATTTC 383
RESULT 11
AAK20973
ID AAK20973 standard; DNA; 401 BP.
AC AAK20973;
XX
XX
XX 05-NOV-2001 (first entry)
DE Human brain expressed single exon probe SEQ ID NO: 20964.
XX
XX Human; brain expressed exon; gene expression analysis; probe;
KW microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;
KW epilepsy; cancer; ss.
XX
XX Homo sapiens.
OS
XX
XX WO200157275-A2.
PN
XX
XX 09-AUG-2001.
PD
XX
XX 30-JAN-2001; 2001WO-US00667.
PF
XX
XX 04-FEB-2000; 2000US-0180312.
PR
XX 26-MAY-2000; 2000US-0207456.
PR
XX 30-JUN-2000; 2000US-0608408.
PR
XX 03-AUG-2000; 2000US-0632366.
PR
XX 21-SEP-2000; 2000US-0234687.
PR
XX 27-SEP-2000; 2000US-0236359.
PR
XX 04-OCT-2000; 2000GB-0024263.
```

```
PR 27-SEP-2000; 2000US-0236359.
PR 04-OCT-2000; 2000GB-0024263.
XX (MOLE-) MOLECULAR DYNAMICS INC.
PA Penn SG, Hanzel DK, Chen W, Rank DR;
XX WPI; 2001-483446/52.
XX Single exon nucleic acid probes for analyzing gene expression in human
PT brains -
XX
PS Example 4; SEQ ID NO: 20964; 650pp + Sequence Listing; English.
CC The present invention provides a number of single exon nucleic acid
CC probes which are derived from genomic sequences expressed in the human
CC brain. They can be used to measure gene expression in brain cell samples,
CC which may enable the diagnosis and improved treatment of nervous system
CC diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia,
CC epilepsy and cancers. The present sequence is one of the probes of the
CC invention.
XX
SQ Sequence 401 BP; 115 A; 92 C; 87 G; 107 T; 0 other;
Query Match 12.8%; Score 159; DB 22; Length 401;
Best Local Similarity 82.1%; Pred. No. 1.7e-34;
Matches 183; Conservative 0; Mismatches 40; Indels 0; Gaps 0;
QY 707 CTGATCTCAAAATTTGAAGAATCTTTTGTCCACCCACACACCCCAAGAAATAATA 766
DB 161 CTGAGCTCTAAATCTGAAGAATCTGTGTCCACCCACACAGCTTCAATTGAAATAAAA 220
QY 767 AACAGGAGGGGAGGATGAAATTTGGCGTCTACCAACCCCTCCAGTAGCAGAAACACCTG 826
DB 221 AACAGGAGGGGAGGATGAAATTTGGCGTCTACCAACCCCTCCAGTAGCAGAAACATCTG 280
QY 827 TACCACTCTCTTCAGTAACAGAAATAGAGACCCCACTGCAAAAGAAATTCGCGGACTGCTA 886
DB 281 TGCCGCTCTCTCGGTAGCAGGAATAGAGACCCCAATACAAAGAAATTTACGCTCTGCTG 340
QY 887 CCATAGCTGGAGAGCCCTTAGGACATTTGCACATTTTCACTATTTC 929
DB 341 CCATAGCTGGAGAGCCCTTAGGACATTTGCACATTTTCACTATTTC 383
RESULT 12
AAK47126
ID AAK47126 standard; DNA; 401 BP.
AC AAK47126;
XX
XX
XX 06-NOV-2001 (first entry)
DE Human bone marrow expressed single exon probe SEQ ID NO: 21683.
XX
XX Human; bone marrow expressed exon; gene expression analysis; probe;
KW microarray; cancer; leukaemia; lymphoma; myeloma; ss.
XX
XX Homo sapiens.
OS
XX
XX WO200157276-A2.
PN
XX
XX 09-AUG-2001.
PD
XX
XX 30-JAN-2001; 2001WO-US00668.
PF
XX
XX 04-FEB-2000; 2000US-0180312.
PR
XX 26-MAY-2000; 2000US-0207456.
PR
XX 30-JUN-2000; 2000US-0608408.
PR
XX 03-AUG-2000; 2000US-0632366.
PR
XX 21-SEP-2000; 2000US-0234687.
PR
XX 27-SEP-2000; 2000US-0236359.
PR
XX 04-OCT-2000; 2000GB-0024263.
```


XX	(MOLE-) MOLECULAR DYNAMICS INC.
PA	Penn SG, Hanzel DK, Chen W, Rank DR;
PI	WFI; 2002-114183/15.
PP	Spatially-addressable set of single exon nucleic acid probes, used to
PT	measure gene expression in human lung samples -
XX	Claim 4; SEQ ID NO 21344; 634pp; English.
PS	The invention relates to a spatially-addressable set of single exon
XX	nucleic acid probes for measuring gene expression in a sample derived
CC	from human lung comprising single exon nucleic acid probes having one of
CC	12614 nucleic acid sequences mentioned in the specification, or their
CC	complements or the 12387 open reading frames derived from the 12614
CC	probes. Also included are a microarray comprising the novel set of
CC	probes; the novel set of probes which hybridize at high stringency to a
CC	nucleic acid expressed in the human lung, comprising (a) contacting the array with
CC	a sample derived from human lung, comprising (a) contacting the array with
CC	a collection of detectably labeled nucleic acids derived from human lung
CC	mRNA, and (b) measuring the label detectably bound to each probe of
CC	the array; identifying exons in a eukaryotic genome, comprising
CC	(a) algorithmically predicting at least one exon from genomic sequences
CC	of the eukaryote; and (b) detecting specific hybridisation of detectably
CC	labeled nucleic acids from eukaryote lung mRNA, to a single exon probe,
CC	having a fragment identical to the predicted exon, the probe is included
CC	in the above mentioned microarray; assigning exons to a single gene,
CC	comprising (a) identifying exons from genomic sequence by the method
CC	above and (b) measuring the expression of each of the exons in several
CC	tissues and/or cell types using hybridisation to a single exon
CC	microarrays having a probe with the exon, where a common pattern of
CC	expression of the exons in the tissues and/or cell types indicates that
CC	the exons should be assigned to a single gene; a peptide comprising one
CC	of 12011 sequences, mentioned in the specification, or encoded by the
CC	probes/open reading frames (ORF). The probes are used for gene
CC	expression analysis, and for identifying exons in a gene, particularly
CC	using human lung derived mRNA and for the study of lung diseases
CC	such as asthma, lung cancer, chronic obstructive pulmonary disease
CC	(COPD), interstitial lung disease (ILD), familial idiopathic pulmonary
CC	fibrosis, neurofibromatosis, tuberous sclerosis, Gaucher's disease,
CC	Niemann-Pick disease, Hermansky-Pudlak syndrome, sarcoidosis, pulmonary
CC	haemochromatosis, pulmonary histiocytosis, lymphangioleiomyomatosis,
CC	pulmonary alveolar proteinosis, Karagener syndrome, fibrocystic
CC	pulmonary dysplasia, primary ciliary dyskinesia, pulmonary hypertension
CC	and hyaline membrane disease. The present sequence is a single exon
CC	probe open reading frame of the invention.
CC	Note: The sequence data for this patent did not form part
CC	of the printed specification, but was obtained in electronic
CC	format directly from WIPO at
XX	ftp.wipo.int/pub/published_pct_sequences.
XX	Sequence 401 BP; 115 A; 92 C; 87 G; 107 T; 0 other;
SQ	Query Match 12.8%; Score 159; DB 24; Length 401;
	Best Local Similarity 82.1%; Pred. No. 1.7e-34;
	Matches 183; Conservative 0; Mismatches 40; Indels 0; Gaps
QY	707 CTGAATCTCAAAATTTTGAAGAATCTTTTGTCACCACCAACCCAAGAAAATAAATA 766
DB	
	161 CTGAGCTCAA AAAATCTGAAGAATCTGTGTCGCCACCCACAGCTTCAATTGAAATAA AA 220
QY	767 AACAGGAGGGGAGGATGAAAAATTTGGCGTCTACACCCCCTCCAGTAGCAGAAACACTTG 826
DB	
	221 AACAGGAGGGGAGGATGAAAAATTTGGCCTATTCTCCAGTGTGCAGAACACTG 280
QY	827 TACCATCTCCTCAGTAACAGAAATAGAGACCCCACTGCAAGAAATTCGCGGACTGCTA 886
DB	
	281 TGCGGCTCCTCTGGTAGCAGGAATAGAGACCCCAATACAAGAAATTTACGCTCTGCTG 340
QY	887 CCATAGCTGGAGGCCCTTAGGACATTTGCATTTTCACTATTC 929

Db 341 CCATAGCTGGAGAGCCCTCAGGACCTTGCTGCTTTCCCTATTTC 383

RESULT 15
ABA60008

ID ABA60008 standard; DNA; 552 BP.

XX AC ABA60008;

XX DT 01-FEB-2002 (first entry)

XX DE Human foetal liver single exon nucleic acid probe #8313.

XX KW Human; foetal liver; gene expression; single exon nucleic acid probe; ss.

XX OS Homo sapiens.

XX PN WO200157277-A2.

XX PD 09-AUG-2001.

XX PF 30-JAN-2001; 2001WO-US00669.

XX PR 04-FEB-2000; 2000US-0180312.

XX PR 26-MAY-2000; 2000US-0207456.

XX PR 30-JUN-2000; 2000US-0608408.

XX PR 03-AUG-2000; 2000US-0632366.

XX PR 21-SEP-2000; 2000US-0234687.

XX PR 27-SEP-2000; 2000US-0236359.

XX PR 04-OCT-2000; 2000GB-0024263.

XX PA (MOLE-) MOLECULAR DYNAMICS INC.

XX PI Penn SG, Hanzel DK, Chen W, Rank DR;

XX DR WPI; 2001-483447/52.

XX PT Human genome-derived single exon nucleic acid probes useful for
PT analyzing gene expression in human fetal liver -

XX PS Claim 1; SEQ ID NO 8313; 639pp + sequence listing; English.

XX CC The invention relates to a single exon nucleic acid probe for
CC measuring human gene expression in a sample derived from human foetal
CC liver. The single exon nucleic acid probes may be used for predicting,
CC measuring and displaying gene expression in samples derived from human
CC foetal liver. The present sequence is a single exon nucleic acid
CC probe of the invention.

CC Note: The sequence data for this patent did not form part of the
CC printed specification, but was obtained in electronic format directly
CC from WIPO at fip.wipo.int/pub/published_pct_sequences.

XX SQ Sequence 552 BP; 164 A; 126 C; 114 G; 148 T; 0 other;

Query Match 12.8%; Score 159; DB 22; Length 552;
Best Local Similarity 82.1%; Pred. No. 2e-34;
Matches 183; Conservative 0; Mismatches 40; Indels 0; Gaps 0

Qy 707 CTGAATCTAAATTTGAAGAATCTTTGTCCACCACCAACACCCAAAGAAATAATA 766
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Db 186 CTGATCTAAATAATCTGAAGAATCTGTTGTCCACCACAGCTTCAATTGAAAAATAAAA 245
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Qy 767 AACAGGAGAGGAGGATGAATAATTGGCGTCTACCCACCCCTCCAGTACGAGAAACACCTG 826
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Db 246 AACAGGAGAGGAGGATGAATAATTGGCGTCTATACGTGCTCTCCAGTTCGAGAAACATCTG 305
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Qy 827 TACCATCTCCTTCAGTAACAGAATAATAGACCCCTACCTGCAAGAATTCGGGGACTGCTA 886
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Db 306 TGCCGCCCTCCTTCGGTAGCAGGATAGACCCCAATACAGAATTTTACGCTCTGCTG 365
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Qy 887 CCATAGCTGGAGAGCCCTTAGGACATTTGACATTTTCACTATTTC 929
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Db 366 CCATAGCTGGAGAGCCCTCAGGACCTTGCTGCTTTTCTATTTC 408
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

Search completed: December 24, 2002, 16:55:30
Job time : 607.404 secs

THIS PAGE BLANK (USPTO)

THIS PAGE BLANK (USPTO)

GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: December 24, 2002, 16:23:56 ; Search time 165.787 Seconds
(without alignments)
13583.723 Million cell updates/sec

Title: US-09-708-724a-3_COPY_1_1000

Perfect score: 1000

Sequence: 1 agccagactaggagtgcgc.....cacacatagatgcagagga 1000

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 2185239 seqs, 1125999159 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : N_Geneseq_101002.*

- 1: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1980.DAT.*
- 2: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1981.DAT.*
- 3: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1982.DAT.*
- 4: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1983.DAT.*
- 5: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1984.DAT.*
- 6: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1985.DAT.*
- 7: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1986.DAT.*
- 8: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1987.DAT.*
- 9: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1988.DAT.*
- 10: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1989.DAT.*
- 11: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1990.DAT.*
- 12: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1991.DAT.*
- 13: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1992.DAT.*
- 14: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1993.DAT.*
- 15: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1994.DAT.*
- 16: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1995.DAT.*
- 17: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1996.DAT.*
- 18: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1997.DAT.*
- 19: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1998.DAT.*
- 20: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1999.DAT.*
- 21: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2000.DAT.*
- 22: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001A.DAT.*
- 23: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001B.DAT.*
- 24: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	ID	Description
1	130	13.0	1620	23	AA565122
2	42	4.2	10732	21	AAA10594
3	36.8	3.7	613	23	AAS86992
4	36.2	3.6	411	24	ABN17146
5	35.2	3.5	3681	24	ABL90758
6	34	3.4	100848	22	AAF28552
7	33.8	3.4	744	20	AA98839
8	33.8	3.4	1597	21	AAA95810
9	33.8	3.4	2168	21	AAA95801

c 10	33.6	3.4	6761	21	AA57362	DNA encoding a hum
c 11	33.4	3.3	637	24	ABN76551	Human ORF1498 cDNA
c 12	33.2	3.3	2543	18	AA74204	Mouse LYST2 cDNA.
c 13	33.2	3.3	16998	24	AD36511	Human Her-1 gene.
c 14	33.2	3.3	197496	24	ABN85584	Human EGFR SEQ ID
c 15	33	3.3	2027	19	AA25979	Human CD33-like pr
c 16	32.6	3.3	1814	22	AAK83065	Human immune/haema
c 17	32.6	3.3	4047	21	AA238853	Human Jurkat cell
c 18	32.6	3.3	38863	21	AA238863	Human Jurkat cell
c 19	32.6	3.3	13646	24	AA220126	Human gene for ret
c 20	32.6	3.3	13646	24	AA220128	Human gene for ret
c 21	32.4	3.2	289	24	ABK73960	Bacillus lichenifo
c 22	32.4	3.2	4273	23	ABL26800	Drosophila melanog
c 23	32.4	3.2	14849	24	ABK12951	DNA encoding mouse
c 24	32.4	3.2	14849	24	ABK24094	Mouse alpha2 macro
c 25	32.4	3.2	29598	19	AAV49654	Human SC2 DNA. Ho
c 26	32.2	3.2	300	23	AA569087	DNA encoding novel
c 27	32.2	3.2	367	22	AA526498	Human cDNA encodin
c 28	32.2	3.2	1107	22	AA040007	Human full length
c 29	32.2	3.2	1832	22	AAK51903	Human polynucleoti
c 30	32.2	3.2	2408	22	AA27634	DNA encoding human
c 31	32.2	3.2	2544	24	ABO91988	Human NF-kB activa
c 32	32.2	3.2	3476	18	AA795898	Novel human gene,
c 33	32.2	3.2	4590	22	AAH24065	Yeast AOD9604-asso
c 34	32	3.2	818	22	AAK82098	Human immune/haema
c 35	31.8	3.2	838	22	AAI95312	Human neuroblastom
c 36	31.8	3.2	2028	23	AA559879	Human novel cytoki
c 37	31.8	3.2	2028	23	ABV22788	Human prostate exp
c 38	31.8	3.2	2028	23	ABV28615	Human prostate exp
c 39	31.8	3.2	6001	24	ABK31214	Signal transductio
c 40	31.8	3.2	15425	22	AA36154	Human cardiovascular
c 41	31.8	3.2	25950	22	AA331518	Human DNA for a no
c 42	31.8	3.2	25950	24	ABO66842	Human polynucleoti
c 43	31.6	3.2	380	24	ABN26293	Human ORFX polynuc
c 44	31.6	3.2	392	22	AAI85371	Human polynucleoti
c 45	31.6	3.2	700	22	AA02740	Human headpin (for

ALIGNMENTS

RESULT 1

AA565122

ID AA565122 standard; cDNA; 1620 BP.

XX AA565122;

XX AA565122;

XX AA565122;

DT 13-FEB-2002 (first entry)

XX DNA encoding novel human diagnostic protein #926.

DE Human; chromosome mapping; gene mapping; gene therapy; forensic;

XX Human; chromosome mapping; gene mapping; gene therapy; forensic;

KW food supplement; medical imaging; diagnostic; genetic disorder; ss.

XX Homo sapiens.

OS Homo sapiens.

XX WO200175067-A2.

PN WO200175067-A2.

XX 11-OCT-2001.

PD 11-OCT-2001.

XX 30-MAR-2001; 2001WO-US08631.

XX 31-MAR-2000; 2000US-0540217.

XX 23-AUG-2000; 2000US-0649167.

PR 23-AUG-2000; 2000US-0649167.

XX (HYSE-) HYSEQ INC.

XX Drmanac RT, Liu C, Tang YT;

XX WPI; 2001-639362/73.

DR P-PSDB; ABG00935.

XX New isolated polynucleotide and encoded polypeptides, useful in

PT

PT diagnostics, forensics, gene mapping, identification of mutations
PT responsible for genetic disorders or other traits and to assess
XX biodiversity -
XX
PS Claim 1; SEQ ID No 926; 103pp; English.
XX
CC The invention relates to isolated polynucleotide (I) and
CC polypeptide (II) sequences. (I) is useful as hybridisation probes,
CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome
CC and gene mapping, and in recombinant production of (II). The
CC polynucleotides are also used in diagnostics as expressed sequence tags
CC for identifying expressed genes. (I) is useful in gene therapy techniques
CC to restore normal activity of (II) or to treat disease states involving
CC (II). (II) is useful for generating antibodies against it, detecting or
CC quantitating a polypeptide in tissue, as molecular weight markers and as
CC a food supplement. (II) and its binding partners are useful in medical
CC imaging of sites expressing (II). (I) and (II) are useful for treating
CC disorders involving aberrant protein expression or biological activity.
CC The polypeptide and polynucleotide sequences have applications in
CC diagnostics, forensics, gene mapping, identification of mutations
CC and to produce other types of data and products dependent on DNA and
CC amino acid sequences. AAS64197-AAS94564 represent novel human
CC diagnostic coding sequences of the invention.
CC Note: The sequence data for this patent did not appear in the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 1620 BP; 408 A; 428 C; 452 G; 332 T; 0 other;

Query Match 13.0%; Score 130; DB 23; Length 1620;
Best Local Similarity 75.7%; Pred. No. 1.6e-29;
Matches 174; Conservative 0; Mismatches 55; Indels 1; Gaps 1;
QY 1 ACCGACTAGACAGTACGACCAAGAGGGGAGGAGTGTGGAGGCACAGGCTGCACCTCT- 59
DB 1177 AGCCACGGGAGGAGTACGAGCAGAGAGAACACAGGCGAGTGGCGGCTGCCTCTC 1236
QY 60 ACTGGTGCCCAAGACCCAGCTGCATGCCAGGCTGCAGTCCAAAGGATACCTCGGTGGG 119
DB 1237 ACTGGTGCCACAGACCTAGCTGTGTTCCAGGCTGCCGTACAGGGGTACTCGGTGG 1296
QY 120 GTCCCTGTCCCAATAGCATCTTAGATCAGCTGCTGAGGCTGGAGCTTCTCCATCCTT 179
DB 1297 GTCCAGGCGCCCTCAGTGGCTGAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1356
QY 180 GAGCATCAGGGGTGTGATCATTTTCCAGGGTTTTCAGACATCCTCGT 229
DB 1357 GAGCAACAGGGGCTGTATCATTTTCCAAAGGATTTTCAGACAAATCCCCGT 1406

RESULT 2
AAAL0594/c
ID AAAL0594 standard; DNA; 10732 BP.
XX
AC AAAL0594;
XX
DT 29-JUN-2000 (first entry)
XX
DE Gene encoding a subunit of cellulose synthase.
XX
KW Cellulose synthase; cellulose production; increase yield; ds.
XX
OS Vigna angularis.
XX
PN JP2000060568-A.
XX
PD 29-FEB-2000.
XX
PF 26-AUG-1998; 98JP-02339998.
XX
PR 26-AUG-1998; 98JP-02339998.
XX

PA (MIZU/) MIZUNO K.
PA (OUIP) OJI PAPER CO.
XX
DR WPI; 2000-342371/30.
DR P-PSDB; AAY85179.
XX
PT A gene encoding a cellulose synthetic equipment - for the improvement
PT in the amount of cellulose synthesised in a plant body
XX
PS Claim 2; Page 14-21; 32pp; Japanese.
XX
CC This sequence represents a gene encoding a subunit of the cellulose
CC synthase complex of Vigna angularis. The invention relates to subunits of
CC cellulose synthetic equipment, that can be used to increase the amount of
CC cellulose synthesised by a plant. The proteins and genes encoding them
CC can also be used to improve the properties of the cellulose being
CC produced by a plant.
XX
SQ Sequence 10732 BP; 3149 A; 1212 C; 2074 G; 2046 T; 2251 other;
Query Match 4.2%; Score 42; DB 21; Length 10732;
Best Local Similarity 14.5%; Pred. No. 0.091;
Matches 82; Conservative 229; Mismatches 247; Indels 8; Gaps 1;
QY 111 TCGGTGGGGTCCCTGTCCCATAGCATCTTAGATCAGCTGCTGAGGCTGGAGCTTCTT 170
DB 9897 YSSRGSDBRGNYNSTNCYDASTDTBYSRCCYTSYSTDSTDTNSTTBSDCYT 9838
QY 171 CCATCTCTTGAGCATCAGGGGTGTGTATCATTTTCAAGGGTTTTCAGACATCCTCGTG 230
DB 9837 TTTBSRSTSDSTSTYRCRSYDATBDSNSTNCCYDASRTBTSTNCYARCTBYDARCS 9778
QY 231 ACCCTGCGAGGGGGCGGTATCATGCGGATCGGTGCTGCTGCTGCTGCTGCTGCTGCTG 290
DB 9777 RDSYSSRGYDANSTSYSSRYSSYSTSYSAKYCAKSTBTBECYDAYDACYDAYDA 9718
QY 291 CCAGCAATCCCATGCCCCACCAATGACATAAATGTTGTGTGGGCGCTTCTTGTGGAAGC 350
DB 9717 NCYSSDSTYTBYSRRCCYDAYSCSYDA-----RCYDACYSYSNSTCYDATBT 9666
QY 351 TCACCTTCTCCTCTGTTGCTCCATCTTCCCAACACAGTACTTCTGCGCATCTCC 410
DB 9665 SRYSTSYSNCYDATTSTRTCTBYSTBTBTTSRCAKCTBDSTAKNSTSYSTTRCTBY 9606
QY 411 TTGTACACCAATGGGAAACTGGTCTCGGAGACTCAGAAACCACTGTGAGCGCTCGA 470
DB 9605 SRSRGYSYCSRSRRCYSCYTDSDSTCYSTYTYAYSCTTSRGYSYDASRSTSYSR 9546
QY 471 GTCTTCCCTGCTCGGCTAACAGGGCATGGAATCAGAGAGAAAGTCACTTCCACCTC 530
DB 9545 CTSTSYSTTDDYSDCYSTTTTBNSTYSDDCTBYSSDRCSRSDSTCNCYSCSDSRY 9486
QY 531 CTGAGGCTGCCAGGCTGAGGCTGGCACACTGAGGCTGACAGGGGCTCTGAAGGCC 590
DB 9485 TTYDACYTYDAKTBCTYTYSDNCCNSTSRCTNSTNSRSTBSRSTCCCTBTTSRGNC 9426
QY 591 AGAGGAGATGGCGGACATAAGGCTGAAGCAACTCTCTGAGCAAGATCTGTTGT 650
DB 9425 YDAYDANSTSYDAYDACYSYDASTBYSYCTBYSNSTYDAYSSRYSCYTCYCDYS 9366
QY 651 GTCTCTCTGAATCTTAGTGGCTTCT 676
DB 9365 STCYTRCAKCTBCNSTSRKSRNTTT 9340
RESULT 3
AAS86992/c
ID AAS86992 standard; cDNA; 613 BP.
XX
AC AAS86992;
XX
DT 13-FEB-2002 (first entry)
XX

DE DNA encoding novel human diagnostic protein #22796.
XX Human; chromosome mapping; gene mapping; gene therapy; forensic;
KW food supplement; medical imaging; diagnostic; genetic disorder; ss.
XX
OS Homo sapiens.
XX WO200175067-A2.
XX 11-OCT-2001.
XX 30-MAR-2001; 2001WO-US08631.
XX 31-MAR-2000; 2000US-0540217.
PR 23-AUG-2000; 2000US-0649167.
XX (HYSE-) HYSEQ INC.
XX Drmanac RT, Liu C, Tang YT;
PI WPI; 2001-639362/73.
DR P-PSDB; ABG22805.
XX
PT New isolated polynucleotide and encoded polypeptides, useful in
PT diagnostics, forensics, gene mapping, identification of mutations
PT responsible for genetic disorders or other traits and to assess
PT biodiversity -
XX
PS Claim 1; SEQ ID No 22796; 103pp; English.
XX
CC The invention relates to isolated polynucleotide (I) and
CC polypeptide (II) sequences. (I) is useful as hybridisation probes,
CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome
CC and gene mapping, and in recombinant production of (II). The
CC polynucleotides are also used in diagnostics as expressed sequence tags
CC for identifying expressed genes. (II) is useful in gene therapy techniques
CC to restore normal activity of (II) or to treat disease states involving
CC (II). (II) is useful for generating antibodies against it, detecting or
CC quantitating a polypeptide in tissue, as molecular weight markers and as
CC a food supplement. (II) and its binding partners are useful in medical
CC imaging of sites expressing (II). (I) and (II) are useful for treating
CC disorders involving aberrant protein expression or biological activity.
CC The polypeptide and polynucleotide sequences have applications in
CC diagnostics, forensics, gene mapping, identification of mutations
CC responsible for genetic disorders or other traits to assess biodiversity
CC and to produce other types of data and products dependent on DNA and
CC amino acid sequences. AAS64197-AAS94564 represent novel human
CC diagnostic coding sequences of the invention.
CC Note: The sequence data for this patent did not appear in the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 613 BP; 151 A; 141 C; 163 G; 137 T; 21 other;

Query Match 3.7%; Score 36.8; DB 23; Length 613;
Best Local Similarity 51.9%; Pred. No. 0.81;
Matches 83; Conservative 0; Mismatches 77; Indels 0; Gaps 0;

Qy 836 GTCCTTGTGCATAGCTCATCCAGCTTGTGTGATACCAATTCAGTCAAGCTGAACAA 895
Db 476 GCCATAGTCCCTTAACCTACACAACTTGCACCAACCACTTTTACGGGGGTTCCTCC 417
Oy 896 GCTGGCAGCTGCTCAACAGGCTTACCAAGACATCATGTTTTTTTTTTTTTTTTCACCAA 955
Db 416 TCTGTCAATATAAACAAGGCTTACCAATTAATCTCTAGTTCTTGTGTGTATCAACCTT 357
Oy 956 ACCTGGACCTGAATGGGGGTGTGGACACACATAGAGTCCA 995
Db 356 AATTAGGTCTGATTGGGTGTGTGCACCAAGGGGCTCCA 317

RESULT 4
ABN17146

ID ABN17146 standard; cDNA; 411 BP.
XX
AC ABN17146;
XX
DT 24-JUN-2002 (first entry)
XX
DE Human ORFX polynucleotide sequence SEQ ID NO:2769.
XX
KW Human; open reading frame; ORFX; gene therapy; cancer; cirrhosis;
KW hyperproliferative disorder; psoriasis; benign tumour; haemorrhage;
KW degenerative disorder; osteoarthritis; neurodegenerative disorder;
KW cardiovascular disease; diabetes mellitus; systemic lupus erythematosus;
KW hypertension; hypothyroidism; cholesterol ester storage disease;
KW immune deficiency; immune disorder; infectious disease;
KW autoimmune disorder; rheumatoid arthritis; autoimmune thyroiditis;
KW myasthenia gravis; gene; ss.
XX
OS Homo sapiens.
XX WO200192523-A2.
XX
PD 06-DEC-2001.
XX
XX 29-MAY-2001; 2001WO-US10836.
XX 30-MAY-2000; 2000US-206132P.
PR 29-AUG-2000; 2000US-228716P.
XX
XX (CURA-) CURAGEN CORP.
XX
XX Shimkets RA, Leach MD;
XX WPI; 2002-106308/14.
XX P-PSDB; ABP01394.
XX
PT Novel human polypeptides and polynucleotides useful for diagnosing,
PT preventing and treating cardiovascular disease, neurodegenerative,
PT hyperproliferative disorders and autoimmune disorders -
XX
PS Disclosure; SEQ ID 2769; 1037pp; English.
XX
CC The present invention describes substantially purified human proteins
CC (referred to as open reading frame, ORFX, where X is 1-11491 (see Table 1
CC in the specification). ABN15762 to ABN27252 encode the human ORFX
CC proteins given in ABP00010 to ABP11500. ORFX proteins are useful for
CC treating or preventing a pathology associated with an ORFX-associated
CC disorder in humans, and in the manufacture of a medicament for treating a
CC syndrome associated with ORFX-associated disorder. ORFX polynucleotide
CC sequences can be used in gene therapy. ORFX sequences can be used in the
CC treatment of cancer, hyperproliferative disorders, cirrhosis of liver,
CC psoriasis, benign tumours, keloid, degenerative disorders, haemorrhage,
CC osteoarthritis, neurodegenerative diseases, disorders related to organ
CC transplantation, cardiovascular diseases, diabetes mellitus, systemic
CC lupus erythematosus, hypertension, hypothyroidism, cholesterol ester
CC storage disease, various immune deficiencies and disorders, infectious
CC diseases, autoimmune disorders such as multiple sclerosis, rheumatoid
CC arthritis, autoimmune thyroiditis, myasthenia gravis, graft-versus-host
CC disease and autoimmune inflammatory eye disease. ORFX proteins are also
CC useful for treating burns, incisions, ulcers, for treating osteoporosis,
CC bone degenerative disorders, or periodontal disease, and for gut
CC protection or regeneration and treatment of lung or liver fibrosis,
CC systemic cytokine damage.
CC N.B. The sequence data for this patent did not form part of the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 411 BP; 79 A; 129 C; 130 G; 73 T; 0 other;

Query Match 3.6%; Score 36.2; DB 24; Length 411;
Best Local Similarity 54.0%; Pred. No. 1;
Matches 74; Conservative 0; Mismatches 63; Indels 0; Gaps 0;

Db 64238 CTGACAGATGTTGCCATGTGCCATGAATGGTAAAGCTTTGGCGACTTTTCATGACGCT 64297
 Qy 618 GAAGCAACCTCTCTGAGCCAAAGATCTGTTGTCTCTCTGAAATCTTAGTGGCTTCTTA 677
 Db 64298 TTACACACCTTGCAGCCAAAGTGAAGATTATGCGCTACCACTGTATGACTGGACAAAGGTC 64357
 Qy 678 AAGCGGGGTGTGATCAGCCATGGGTATCAGACACTGGAGTCCAGTAGCTGCTAGGTGG 737
 Db 64358 AAAGAGCGTGAACCGGCTATATGCTTACCGATGAGGCTCTCTCATGAACGATATCTGG 64417
 Qy 738 GACACGGGCAACATTCCTACTGTCAGACCACTGTCACGGAGTG 779
 Db 64418 ACAGCCTATGCCAATTTACCTGATGACTTGGCCAAAGGCTTG 64459

RESULT 7
 AAX98839
 ID AAX98839 standard; cDNA; 744 BP.
 AC AAX98839;
 XX
 DT 24-SEP-1999 (first entry)
 XX
 DE Human validated cancer cell derived cDNA #161.
 XX
 KW Cancer; human; colon; breast; lung; transmembrane receptor; ATPase;
 KW integral membrane protein; aspartyl protease; GATA family; wnt family;
 KW transcription factor; G-protein alpha subunit; protein phosphatase;
 KW phospholipase; protein binding; diacylglycerol binding protein; trypsin;
 KW protein kinase; tyrosine phosphatase; developmental signalling protein;
 KW WW/rsp5/WMP domain; therapy; forensic; genetic mapping; diagnostic;
 KW detection; treatment; cervical; melanoma; colorectal adenocarcinoma;
 KW Wilm's tumour; retinoblastoma; sarcoma; myosarcoma; lung carcinoma;
 KW leukemia; lymphoma; dysplasia; hyperplasia; endometrium; adrenal;
 KW prostate; ss.
 XX
 OS Homo sapiens.
 XX
 PN W09933982-A2.
 XX
 PD 08-JUL-1999.
 XX
 XX 22-DEC-1998; 98WO-US27610.
 XX
 XX 21-DEC-1998; 98US-0217471.
 PR 23-DEC-1997; 97US-0068755.
 PR 03-APR-1998; 98US-0080664.
 PR 21-OCT-1998; 98US-0105234.
 PR 27-OCT-1998; 98US-0105877.
 XX
 PA (CHIR) CHIRON CORP.
 PA (HYSE-) HYSEQ INC.
 XX
 PI Crkvenjakov R, Dickson M, Drmanac R, Drmanac S;
 PI Escobedo J, Garcia PD, Garcia V, Giese K, Innis MA;
 PI Jones LW, Kassam A, Kennedy GC, Kita D, Labat I;
 PI Lamson G, Leshkowitz D, Pot D, Randazzo F, Reinhard C;
 PI Stache-Crain B, Sudduth-Klinger J, Williams LJ;
 XX
 DR WPI; 1999-430243/36.
 XX
 XX New isolated human polynucleotides
 XX
 XX Claim 1; Page 478; 591pp; English.
 XX
 CC This invention describes novel isolated human polynucleotides obtained
 CC by screening for differential expression in colon cancer, breast cancer
 CC and lung cancer cell lines. The polynucleotides of the invention are
 CC represented in AAX98275-X99118 and encode polypeptides of protein
 CC families selected from 4 transmembrane segments integral membrane
 CC proteins, 7 transmembrane receptors, ATPases associated with various
 CC cellular activities (AAA), eukaryotic aspartyl proteases, GATA family of
 CC transcription factors, G-protein alpha subunit, phospholipase or

CC diacylglycerol binding proteins, protein kinase, protein phosphatase 2C,
 CC protein tyrosine phosphatase, trypsin, wnt family of developmental
 CC signalling proteins and WW/rsp5/WMP domain containing proteins. The
 CC encoded polypeptides also have a functional domain selected from Ank
 CC repeat, basic region plus leucine zipper transcription factors,
 CC bromodomain, EF-hand, SH3 domain, WD domain/G-beta repeats, zinc finger
 CC (C2H2 type), zinc finger (CCHC class), and zinc-binding metalloprotease
 CC domain. The polynucleotides encode polypeptides with similarity to known
 CC protein families and are predicted to have similar properties. The novel
 CC polynucleotides can be used to develop products for use as therapeutic
 CC agents and in forensics, genetic analysis, mapping and diagnostic
 CC applications. In particular, the product can be used for the detection
 CC and management of cancers. They can be used for treating e.g. cervical
 CC cancers, melanomas, colorectal adenocarcinomas, Wilm's tumour, sarcomas,
 CC retinoblastoma, myosarcomas, lung carcinomas, leukemias, such as chronic
 CC myelogenous leukemia, promyelocytic leukemia, monocytic leukemia, and
 CC myeloid leukemia, and lymphomas such as histiocytic lymphoma, anhydric
 CC hereditary ectodermal dysplasia, congenital alveolar dysplasia,
 CC epithelial dysplasia of the cervix, fibrous dysplasia of bone, and
 CC mammary dysplasia, hyperplasias, e.g. endometrial, adrenal, breast,
 CC prostate or thyroid hyperplasias or pseudoepitheliomatous hyperplasia of
 CC the skin.
 XX

SQ Sequence 744 BP; 189 A; 169 C; 188 G; 173 T; 25 other;

Query Match 3.4%; Score 33.8; DB 20; Length 744;
 Best Local Similarity 50.08; Pred. No. 7.7;
 Matches 110; Conservative 0; Mismatches 108; Indels 2; Gaps 1;
 Qy 736 GGGACACGGGCAACATTCCTACTTCAGACCACTGCAGGAGTGGATAAAGAGAGAGTTC 795
 Db 202 GGGCCACTGGGGCATTTTNCACGTTAAACAGCAGCTGCCACTGGCAAAGAGTACTCGCC 261
 Qy 796 TGTGTGGGAATCTCCTTTGTTGGTATCATCAGGAGGTGAAGTCTTTGTTCATAGCCTATA 855
 Db 262 AATGTTGGCATCTCAGATGTGGGCCCCAGGAGTCTGGGAGCTACTTTGAACAG--GGCTA 319
 Qy 856 TCCAGCTTGTGTGATACCAATTCAGTGAAGCTGGCAAGCTGCTCAACACAGG 915
 Db 320 TCCATTTCATTGTCACCAACCAAGGCTATGAGCCACCACCATGCTGCTGGAGTAGTCAAG 379
 Qy 916 CCTACCAAGACATCATGTTTTTTTTTTTTTTTTTTTTTCCACCAA 955
 Db 380 GGAATAAGACACTCTCCTTGTCTTGTGTTAACTCAATCAA 419

RESULT 8
 AAA95810/c
 ID AAA95810 standard; cDNA; 1597 BP.
 XX
 AC AAA95810;
 XX
 DT 09-MAR-2001 (first entry)
 XX
 DE Tobacco cDNA clone T3.
 XX
 KW Tobacco; T3; MAR binding filament-like protein 1; MFPI;
 KW matrix attachment region; MAR; NCMFPI-2; anchor protein; ss.
 XX
 OS Nicotiana tabacum.
 XX
 PN W0200061615-A2.
 XX
 PD 19-OCT-2000.
 XX
 XX 12-APR-2000; 2000WO-US09723.
 XX
 PR 12-APR-1999; 99US-0128900.
 XX
 PA (DUPO) DU PONT DE NEMOURS & CO E I.
 XX
 PI Harder PA, Meier I;
 XX

Db 64 GTGGAGCCCCGACCCAGCGCTGCTTCCTGAGCGTGGGT 26

RESULT 12
AAT74204/c
ID AAT74204 standard; cDNA; 2543 BP.

XX
AC AAT74204;

XX 10-FEB-1998 (first entry)

XX Mouse LYST2 cDNA.

XX LYST2; mouse; lysosomal trafficking regulator;

KW Chediak-Higashi syndrome; CH syndrome; autoimmune disease; tumour;

KW Alzheimer's disease; motor neuron disease; Parkinson's disease;

KW acute tubular necrosis; glomerulonephritis; glomerulosclerosis;

KW vaccine; therapy; diagnosis; ss.

XX

OS Mus musculus.

XX

XX Key Location/Qualifiers

FT CDS 3..2114

FT /*tag= a

XX

PN WO9728262-A1.

XX

PD 07-AUG-1997.

XX

PF 31-JAN-1997; 97WO-US01748.

XX

PR 23-DEC-1996; 96US-0034346.

XX

PR 01-FEB-1996; 96US-0011146.

XX

PR 20-DEC-1996; 96US-0033599.

XX

PA (UYFL) UNIV FLORIDA.

XX

PI Barbosa-Alleyne MDFS, Kingsmore SF;

XX

DR WPI; 1997-402616/37.

XX

DR P-PSDB; AAW23599.

XX

XX Mammalian lysosomal trafficking regulators LYST1, LYST1, LYST2 and

PT LYST2 - useful to diagnose Chediak-Higashi syndrome

XX

PS Claim 7; Page 113-114; 237pp; English.

XX

CC This mouse LYST2 (lysosomal trafficking regulator) cDNA sequence

CC was isolated from a mouse embryo cDNA library using a probe

CC corresponding to human LYST2 (see AAT74203). Murine LYST2 nucleic acids

CC can be used in methods for the recombinant expression of LYST2

CC polypeptides (see AAW23599) useful in various pharmacological and

CC immunological applications, as well as in methods for detecting

CC LYST2 genes and gene mutations.

XX

SQ Sequence 2543 BP; 658 A; 637 C; 615 G; 633 T; 0 other;

Query Match 3.3%; Score 33.2; DB 18; Length 2543;

Best Local Similarity 50.6%; Pred. No. 23;

Matches 80; Conservative 0; Mismatches 78; Indels 0; Gaps 0;

QY 331 GRGGGCTCTTTCTGGAAGCTCACCTTCCTCCTCTTTTGGCCCTCCATCTCCCAACC 390

Db 1358 GTGGGCAATTGATCTGGTCTACAGGCTGTGTGATCTGCCGCTTGTTCACACC 1299

QY 391 AGTACTTCTGGGCAATCCTCCTGTGTACACCAATGGGAAACCTGGGCTGGAGACTCAGA 450

Db 1298 AAGAGTATTTCGGCATTAATGGGTCCATCTCAATGGGAAGATGTGTGTATCCAAAGGA 1239

QY 451 AACCACTGTGGAGGCTCGAGTCTCCCTCTCCCTGCTCCCTGGC 488

Db 1238 GTATCGGGGGCTCTCTCTGAGGCCGACTGTGTGTGCC 1201

RESULT 13
AAD36511
ID AAD36511 standard; DNA; 169998 BP.

XX
AC AAD36511;

XX 09-AUG-2002 (first entry)

XX Human Her-1 gene.

XX Human; epidermal growth factor receptor; hyperproliferative disease;

KW Her1; prophylaxis; psoriasis; tumour; cancer; gene; ds.

XX Homo sapiens.

XX Key Location/Qualifiers

FT exon 1208..1472

FT /*tag= a

FT intron 1473..124390

FT /*tag= b

FT exon 124391..124544

FT /*tag= c

FT intron 124545..125409

FT /*tag= d

FT exon 125410..125595

FT /*tag= e

FT intron 125596..128711

FT /*tag= f

FT exon 128712..128848

FT /*tag= g

FT intron 128849..133400

FT /*tag= h

FT exon 133401..133469

FT /*tag= i

FT intron 133470..134652

FT /*tag= j

FT exon 134653..134773

FT /*tag= k

FT intron 134774..136116

FT /*tag= l

FT exon 136117..136261

FT /*tag= m

FT intron 136262..137936

FT /*tag= n

FT exon 137937..138053

FT /*tag= o

FT intron 138054..138637

FT /*tag= p

FT exon 138638..138766

FT /*tag= q

FT intron 138767..138864

FT /*tag= r

FT exon 138865..138940

FT /*tag= s

FT intron 138941..139765

FT /*tag= t

FT exon 139766..139860

FT /*tag= u

FT intron 139861..142245

FT /*tag= v

FT exon 142246..142445

FT /*tag= w

FT intron 142446..143605

FT /*tag= x

FT exon 143606..143738

FT /*tag= y

FT intron 143739..145838

FT /*tag= z

FT exon 145839..145931

FT /*tag= aa

FT intron 145932..147385

FT /*tag= ab

```
FT exon 147386..147544
FT /tag= ac
FT intron 147545..153274
FT /tag= ad
FT exon 153275..153321
FT /tag= ae
FT intron 153322..155088
FT /tag= af
FT exon 155089..155231
FT /tag= ag
FT intron 155232..156025
FT /tag= ah
FT exon 156026..156151
FT /tag= ai
FT intron 156152..156826
FT /tag= aj
FT exon 156827..156928
FT /tag= ak
FT intron 156929..163399
FT /tag= al
FT exon 163400..163586
FT /tag= am
XX WO200226758-A1.
PN XX
XX 04-APR-2002.
XX
XX 28-SEP-2001; 2001WO-US30551.
XX
XX 29-SEP-2000; 2000US-0676610.
XX
XX (ISIS-) ISIS PHARM INC.
XX
XX Bennett CF, Wyatt JR, Freier SM;
XX WPI; 2002-394234/42.
XX
XX Novel antisense oligonucleotide that specifically hybridizes with and
XX inhibits nucleic acid encoding epidermal growth factor receptor, useful
XX for treating hyperproliferative disease such as cancer or psoriasis -
XX Example 19; Page 67-121; 169pp; English.
XX
XX The invention relates to an antisense oligonucleotide targetted to a
XX nucleic acid molecule encoding human epidermal growth factor receptor
XX (Her1) to inhibit its expression. The antisense compounds are useful
XX for treating diseases or conditions associated with Her-1 such as
XX hyperproliferative diseases especially cancer (lung, ovarian, colon
XX or prostate cancer) and psoriasis. They are also useful as research
XX reagents, diagnostics, therapeutics, kits and prophylactically e.g.
XX to prevent or delay tumour formation. The present sequence is
XX human Her-1 gene.
XX
XX Sequence 169998 BP; 46143 A; 38164 C; 37751 G; 47940 T; 0 other;
XX
XX Query Match 3.3%; Score 33.2; DB 24; Length 169998;
XX Best Local Similarity 59.6%; Pred. No. 2.1e+02;
XX Matches 56; Conservative 0; Mismatches 38; Indels 0; Gaps 0;
XX
XX Qy 506 AGAGAGAAAGTCATCTTCCACCTCCTGAAGGCTGCCAGCGTCAGGGCTTGGCACACTGA 565
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX Db 57186 AGGTTGTGTAGGTAGCCAGACCATCATCTAGTGGGCGACGCCAGGCGGCGGCACCGT 57245
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX Qy 566 GGCTGACAGGGCGCTCTCTGAAGGCCAGAGGAGAT 599
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX Db 57246 TGCAGCCTGCATCTCTTCTAAAGGGCAGAGCAAT 57279
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX
XX RESULT 14
XX ABN85584
XX ID ABN85584 standard; DNA; 197496 BP.
XX
XX AC ABN85584;
```

```
XX 09-SEP-2002 (first entry)
XX
XX Human EGFR SEQ ID NO 10.
XX
XX Human; EGFR; HER2-neu; chemotherapeutic regimen; tumour; cancer;
XX receptor tyrosine kinase; epidermal growth factor receptor;
XX gene expression; ds.
XX
XX Homo sapiens.
XX
XX WO200244413-A2.
XX
XX 06-JUN-2002.
XX
XX 09-NOV-2001; 2001WO-US43035.
XX
XX 01-DEC-2000; 2000US-250122P.
XX
XX 04-DEC-2000; 2000US-250469P.
XX
XX 11-JUN-2001; 2001US-0877177.
XX
XX (RESP-) RESPONSE GENETICS INC.
XX
XX Danenberg KD;
XX WPI; 2002-537460/57.
XX
XX Determining chemotherapeutic regimen of receptor tyrosine kinase
XX targeted agent for treating tumor by examining EGFR and/or HER2-neu
XX mRNA amount in tumor cells, comparing it to predetermined threshold
XX expression level -
XX
XX Disclosure; Page 71-124; 125pp; English.
XX
XX The invention relates to determining the chemotherapeutic regimen of
XX receptor tyrosine kinase targeted agent for treating tumour by amplifying
XX mRNA from tumour and non-malignant tissues using a primer pair that
XX hybridises to epidermal growth factor receptor (EGFR) and/or HER2-neu
XX gene (1), quantitating and obtaining differential expression levels of
XX amplified mRNA and comparing the differential expression levels and
XX threshold levels for expression of (1). The method is useful for
XX assessment of clinical treatment of a patient and as a diagnostic or
XX prognostic tool for a range of cancers including breast, head and neck,
XX lung, oesophageal and colorectal cancer. The present sequence is that of
XX the human EGFR DNA sequence used in methods of the invention.
XX
XX Sequence 197496 BP; 53640 A; 44528 C; 43228 G; 56100 T; 0 other;
XX
XX Query Match 3.3%; Score 33.2; DB 24; Length 197496;
XX Best Local Similarity 59.6%; Pred. No. 2.3e+02;
XX Matches 56; Conservative 0; Mismatches 38; Indels 0; Gaps 0;
XX
XX Qy 506 AGAGAGAAAGTCATCTTCCACCTCCTGAAGGCTGCCAGCGTCAGGGCTTGGCACACTGA 565
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX Db 65186 AGGTTGTGTAGGTAGCCAGACCATCATCTAGTGGGCGACGCCAGGCGGCGGCACCGT 65245
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX Qy 566 GGCTGACAGGGCGCTCTCTGAAGGCCAGAGGAGAT 599
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX Db 65246 TGCAGCCTGCATCTCTTCTAAAGGGCAGAGCAAT 65279
XX ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
XX
XX RESULT 15
XX AAV25979/c
XX ID AAV25979 standard; cDNA; 2027 BP.
XX
XX AC AAV25979;
XX
XX 21-JUL-1998 (first entry)
XX
XX Human CD33-like protein encoding cDNA.
XX
XX Human; CD33; CD33-like protein; tumour; inflammatory disease;
XX leukaemia; bone marrow; monocyte; haematopoietic; antibody; ss.
```

```
XX OS Homo sapiens.
XX FH Key
XX CDS Location/Qualifiers
XX     37..1692
XX     /*tag= a
XX     /product= "CD33-like protein"
XX sig_peptide
XX     37..81
XX     /*tag= b
XX
XX W09806733-A1.
XX
XX PD 19-FEB-1998.
XX
XX PF 09-AUG-1996; 96WO-US13007.
XX
XX PR 09-AUG-1996; 96WO-US13007.
XX
XX PA (HUMA-) HUMAN GENOME SCI INC.
XX
XX PI Gentz RL, Ni J, Rosen CA;
XX
XX WPI; 1998-159451/14.
XX P-PSDB; AAW55884.
XX
XX New nucleic acid nearly identical to sequence encoding CD33-like
XX protein - useful in, e.g. diagnosis of tumour or inflammatory
XX disease and purging bone marrow monocytic haematopoietic cells from
XX leukaemia patients
XX
XX Claim 1; Fig 1; 83pp; English.
XX
XX The present sequence encodes a CD33-like protein. The present sequence
XX was obtained by sequencing the HMOC14 cDNA clone contained in ATCC
XX Deposit No. 97521. An isolated antibody that binds specifically to the
XX CD33-like protein may be used for the detection of the CD33-like protein
XX or its mRNA, and so is useful for, e.g. diagnosing a tumour or
XX inflammatory disease. The antibody (especially an immunotoxin), can
XX also be used to remove or deplete haematopoietic cells expressing the
XX CD33-like protein antigen, which can be used to purge bone marrow
XX monocytic haematopoietic cells obtained from a leukaemia patient, which
XX can subsequently be reinfused into a patient previously subjected to
XX myeloablative chemotherapy. The antibody can also be used as an
XX antagonist to inhibit the CD33-like protein receptor signalling pathway,
XX useful for inhibiting the growth or selective killing of tumour cells.
XX
XX Sequence 2027 BP; 477 A; 602 C; 547 G; 401 T; 0 Other;
XX
Query Match 3.3%; Score 33; DB 19; Length 2027;
Best Local Similarity 53.5%; Pred. NO. 23;
Matches 69; Conservative 0; Mismatches 60; Indels 0; Gaps 0;
QY 530 CCTGAGGCTGCCAGGCTGGCAGCTGACACTGAGGCTGACAGGGGCTTCTGAAGGC 589
Db | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
1960 CATGATGATGCTCTATTACAGATGGGAATGGAACATGAGGCACAGAGGGATGTCAGTGAATTG 1901
QY 590 CAGAGGAGATGCCGGGACATAAGGCTGAAGCAACCTCTCTGAGCCAAAGATCTGTTTG 649
Db | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
1900 CTTGATGACAGTGCCAGGSCCTAGATTGAGCCCACTCTCTAATTCATCATCTGTTTT 1841
QY 650 TGTCCTCCT 658
Db | | | |
1840 CTTTCTGCT 1832
```

Search completed: December 24, 2002, 16:59:46
Job time : 421.786 secs

GenCore version 5.1.3
Copyright (c) 1993 - 2002 CompuGen Ltd.

OM nucleic - nucleic search, using sw model

Run on: December 24, 2002, 16:23:56 ; Search time 165.952 Seconds
(without alignments)
13583.723 Million cell updates/sec

Title: US-09-708-724A-3_COPY_70000_71000

Perfect score: 1001

Sequence: 1 ggagatggataaacctgtg.....ccattcaggagtctatgtg 1001

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 2185239 seqs, 112599159 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : N_Geneseq_101002.*

- 1: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1980.DAT.*
- 2: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1981.DAT.*
- 3: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1982.DAT.*
- 4: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1983.DAT.*
- 5: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1984.DAT.*
- 6: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1985.DAT.*
- 7: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1986.DAT.*
- 8: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1987.DAT.*
- 9: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1988.DAT.*
- 10: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1989.DAT.*
- 11: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1990.DAT.*
- 12: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1991.DAT.*
- 13: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1992.DAT.*
- 14: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1993.DAT.*
- 15: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1994.DAT.*
- 16: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1995.DAT.*
- 17: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1996.DAT.*
- 18: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1997.DAT.*
- 19: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1998.DAT.*
- 20: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1999.DAT.*
- 21: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2000.DAT.*
- 22: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001A.DAT.*
- 23: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001B.DAT.*
- 24: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1001	100.0	2578	21	AAZ93065
2	584.6	58.4	4977	22	HERV-AVL3-B tumour
3	455.6	45.5	2202	21	Human genomic DNA
4	454	45.4	2166	22	Human secreted pro
5	418.6	41.8	808	22	Human CDNA sequenc
6	217.8	21.8	1537	24	Human CDNA clone (
7	192.6	19.2	4823	24	Human neurogenesis
8	180.4	18.0	189	22	Human prostate spe
9	180.4	18.0	189	22	Human breast cancer
					Human breast cancer

10	178.8	17.9	184	22	AA119385	Human breast cancer
11	174.4	17.4	176	22	AA126090	Human breast cancer
12	171.8	17.2	200	22	AA108998	Human breast cancer
13	164.8	16.5	273	24	AB180406	Human ovarian cancer
14	152.8	15.3	660	23	AA92505	DNA encoding novel
15	152.6	15.2	2652	22	AAH15033	Human cDNA sequenc
16	151.8	15.2	522	22	AA143362	Probe #12048 used
17	151.8	15.2	1384	24	AA144826	Human cancer cell
18	151.8	15.2	1384	24	AA144827	Human cancer cell
19	151.4	15.1	609	24	ABQ56844	Human colon cancer
20	151	15.1	1715	22	AA541104	CDNA encoding nove
21	149.4	14.9	1773	23	AA92506	DNA encoding novel
22	148	14.8	932	23	AA569023	DNA encoding novel
23	131.4	13.1	443	22	AA119260	Human breast cancer
24	129.4	12.9	131	22	AA126310	Human breast cancer
25	124.4	12.4	572	22	ABA61773	Human foetal liver
26	124.4	12.4	572	22	AAK10083	Human brain expres
27	124.4	12.4	572	22	AAK35978	Human bone marrow
28	124.4	12.4	572	22	AA141692	Probe #10378 used
29	121.4	12.1	625	22	AA108565	Human breast cancer
30	103.2	10.3	254	20	AA127261	Prostate-tumour de
31	103.2	10.3	465	22	ABA57306	Human foetal liver
32	103.2	10.3	465	22	ABA36859	Probe #5325 for ge
33	103.2	10.3	465	22	AAK05341	Human brain expres
34	103.2	10.3	465	22	AAK30934	Human bone marrow
35	103.2	10.3	465	22	AA115470	Probe #5403 for ge
36	103.2	10.3	465	22	AA136847	Probe #5533 used t
37	103.2	10.3	465	24	ABS05682	Human genome-deriv
38	103.2	10.3	51402	22	AAK72363	Human immune/haema
39	100.6	10.0	143068	21	AAK721105	Human low adenosin
40	100.6	10.0	143068	21	AAK721272	Human low adenosin
41	100.6	10.0	143068	21	AAA34983	Human adenosine re
42	100.6	10.0	143068	21	AAA35150	Human adenosine re
43	100.6	10.0	143068	24	ABL68124	Ovary cancer relat
44	100.6	10.0	149412	21	AAA35151	Human adenosine re
45	100.6	10.0	152740	21	AAK71273	Human low adenosin

ALIGNMENTS

RESULT 1	
AAZ93065	
ID	AAZ93065 standard; DNA; 2578 BP.
XX	
AC	AAZ93065;
XX	
DT	19-JUN-2000 (first entry)
XX	
DE	HERV-AVL3-B tumour associated polypeptide coding sequence.
XX	
KW	Tumour; tumour associated antigen; retrovirus; antisense;
KW	treatment; probe; primer; HLA; cytotoxic T-lymphocyte; cancer;
KW	testis; antibody; ss.
XX	
OS	Homo sapiens.
XX	
PH	Key
LTR	Location/Qualifiers
FT	82..148
FT	/*tag= a
FT	/label= 5' LTR region
FT	2197..2480
FT	/*tag= b
FT	/label= 3' LTR region
XX	
PN	WO200006598-A1.
XX	
PD	10-FEB-2000.
XX	
PF	15-JUL-1999; 99WO-US16236.
XX	
PR	29-JUL-1998; 98US-O124398.
XX	

PA (LUDW-) LUDWIG INST CANCER RES.
XX
PI Coulie P, Boon-falleur T;
XX
DR WPT: 2000-205453/18.
DR P-FSDB: AAY82952.
XX
PT Novel nucleic acids encoding melanoma associated gene products and
PT their fragments and variants, useful for treating endogenous retrovirus
PT mediated tumors, especially melanomas
XX
PS Claim 1; Figure 3; 77pp; English.
XX
CC Tumor associated disorders (e.g. endogenous retrovirus mediated
CC tumors, especially melanomas) can be treated or ameliorated by
CC administering antisenese nucleic acid to reduce the expression of
CC tumour associated genes such as HERV-AVL3-B. Progression of
CC a disorder characterized by the expression of the HERV-AVL3-B
CC endogenous retrovirus tumor rejection antigen (ERTRA) can be
CC diagnosed or monitored by contacting a non-testis biological
CC sample with an agent that binds to the complex and determining
CC the interaction. A disorder can also be treated by administering
CC an agent that enriches the presence of HLA and HERV-AVL3-B ERTRA
CC or by administering autologous cytotoxic T-cells sufficient to
CC ameliorate the disorder. Fragments of the HERV-AVL3-B coding sequence
CC are useful as probes or amplification primers for determining the
CC expression of HERV-AVL3-B genes, to express tumor associated
CC polypeptides in vivo and in vitro and to prepare fragments of such
CC polypeptides to synthesize antibodies. Antigenic peptides of
CC HERV-AVL3-B can be useful for generating antibodies either alone or
CC as fusion proteins, as components of immunoassay and for determining
CC the binding specificity of HLA molecules and/or cytotoxic T
CC lymphocyte (CTL) for HERV-AVL3-B proteins.
XX
SQ Sequence 2578 BP; 785 A; 573 C; 515 G; 705 T; 0 other;

Query Match 100.0%; Score 1001; DB 21; Length 2578;
Best Local Similarity 100.0%; Pred. No. 4.le-313;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GGAGATGATAACCGTGTGAGTGGCCCTCAAGTCTGTGGCAGCATGGAATGGGAGCTG 60
DB 259 GGAGATGATAACCGTGTGAGTGGCCCTCAAGTCTGTGGCAGCATGGAATGGGAGCTG 318
QY 61 GAGGATACATGGATCCCAACTACAGGCCAGCTCCTCCAGTATGAGCCATGAGCCAGT 120
DB 319 GAGGATACATGGATCCCAACTACAGGCCAGCTCCTCCAGTATGAGCCATGAGCCAGT 378
QY 121 GAATCTGAATGTGAAGATGGAATGAAGACCGACGAGAGTCACTGACGTCACACCTCAT 180
DB 379 GAATCTGAATGTGAAGATGGAATGAAGACCGACGAGAGTCACTGACGTCACACCTCAT 438
QY 181 ACATCGGGTCAAGTCAAGAAACACACACAGCAAGCTGAGAATCTGTGTAGTCCAGGG 240
DB 439 ACATCGGGTCAAGTCAAGAAACACACACAGCAAGCTGAGAATCTGTGTAGTCCAGGG 498
QY 241 TCAGCAAAAAACCCCTGACTCCATGTTTATGGCCATGCTAGCTGAATATCTGTGCAGT 300
DB 499 TCAGCAAAAAACCCCTGACTCCATGTTTATGGCCATGCTAGCTGAATATCTGTGCAGT 558
QY 301 ATGATTTTCTGTGAGAGCAAAAAACATATTTGGGCATATTTTCCCTAACCCACCGGTAGT 360
DB 559 ATGATTTTCTGTGAGAGCAAAAAACATATTTGGGCATATTTTCCCTAACCCACCGGTAGT 618
QY 361 GTGATCATACTCTGAGCAGCACTCCTCTGAGATATATCATGATCAGGAGCATCAGTA 420
DB 619 GTGATCATACTCTGAGCAGCACTCCTCTGAGATATATCATGATCAGGAGCATCAGTA 678
QY 421 CCAGGACCTCTAACTCCCCCTGACACAGCAAGTAATAGACTCTCAATAACAATGGTATCAAT 480
DB 679 CCAGGACCTCTAACTCCCCCTGACACAGCAAGTAATAGACTCTCAATAACAATGGTATCAAT 738
QY 481 TATACCACTCCATTGGAGGAGTTCCTTTATGTGTACCCAGGATACATTTGCTCAACTGC 540

DB 739 TATACCACTCCATTGGAGGAGCTTCCTTTATGTGTACCCAGGATACATTTCTCAACTGC 798
QY 541 AGTTGCCCTTGAGTTTATGATCCCAAGCATGGTTGAGTTTACCATAAAAAAATTTATGTACTA 600
DB 799 AGTTGCCCTTGAGTTTATGATCCCAAGCATGGTTGAGTTTACCATAAAAAAATTTATGTACTA 858
QY 601 TTAGACCTTAGCTTTTATTAATATTACTTTGTGTAGTTACTAATCACTCTCGGCCCATCAC 660
DB 859 TTAGACCTTAGCTTTTATTAATATTACTTTGTGTAGTTACTAATCACTCTCGGCCCATCAC 918
QY 661 CCAATTTGCTACTGATTATACAGATGGCTCCCTTTGATAATTTCTACACCCCTCTCTGG 720
DB 919 CCAATTTGCTACTGATTATACAGATGGCTCCCTTTGATAATTTCTACACCCCTCTCTGG 978
QY 721 GCCCACTCTCTTGGCCCTTAGCTAGACAATAGTCCATGTTAATGGGAGACATTTATTGAC 780
DB 979 GCCCACTCTCTTGGCCCTTAGCTAGACAATAGTCCATGTTAATGGGAGACATTTATTGAC 1038
QY 781 TGGGTCCTCTGTGTCATTAAGATGGGAGAGATGAGATGAGATGAGATGAGATGAGATGAGAT 840
DB 1039 TGGGTCCTCTGTGTCATTAAGATGGGAGAGATGAGATGAGATGAGATGAGATGAGATGAGAT 1098
QY 841 CACTGGCACTGGTGGCGAACTTTAAACATCTCTTCACTTCAACACACTGGGATTCATCC 900
DB 1099 CACTGGCACTGGTGGCGAACTTTAAACATCTCTTCACTTCAACACACTGGGATTCATCC 1158
QY 901 CAATCTGCATGCAACTTGTCTTGGCATGGAACGGCTTTAGCCCACTTTGCCTCAATGG 960
DB 1159 CAATCTGCATGCAACTTGTCTTGGCATGGAACGGCTTTAGCCCACTTTGCCTCAATGG 1218
QY 961 CATTATCAAGAAAGAGAGAGTCCCAATTCAGGAGTCTATGTG 1001
DB 1219 CATTATCAAGAAAGAGAGAGTCCCAATTCAGGAGTCTATGTG 1259

RESULT 2
AAS26628
ID AAS26628 standard; DNA; 4977 BP.
XX AC AAS26628;
XX AC AAS26628;
DT 07-NOV-2001 (first entry)
XX
DE Human genomic DNA encoding partial novel secreted protein, Seq ID 1602.
KW Human; immunosuppressive; antiarthritic; ds; antirheumatic;
KW cytostatic; cardiant; vasotropic; cerebroprotective; nootropic;
KW neuroprotective; antibacterial; virucide; fungicide; ophthalmological;
KW vulnary; secreted protein; rheumatoid arthritis;
KW hyperproliferative disorder; cardiovascular disorder; cardiac arrest;
KW cerebrovascular disorder; cerebral ischaemia; angiogenesis;
KW nervous system disorder; Alzheimer's disease; infection; ocular disorder;
KW corneal infection; wound healing; epithelial cell proliferation;
KW skin ageing; food additive; preservative; antiproliferative.
XX Homo sapiens.
XX
XX WO200155322-A2.
PN 02-AUG-2001.
XX
PD 17-JAN-2001; 2001WO-US01341.
XX
PR 31-JAN-2000; 2000US-0179065.
PR 04-FEB-2000; 2000US-0180628.
PR 24-FEB-2000; 2000US-0184664.
PR 02-MAR-2000; 2000US-0186350.
PR 16-MAR-2000; 2000US-0189874.
PR 17-MAR-2000; 2000US-0190076.
PR 18-APR-2000; 2000US-0198123.
PR 19-MAY-2000; 2000US-0205515.
PR 07-JUN-2000; 2000US-0209467.

be used in alleviating symptoms associated with the disorders and in diagnostic immunoassays e.g. radioimmunoassays or enzyme linked immunosorbant assays (ELISA). Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection, and many other disorders listed in the specification. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities, fat content, lipid, protein, carbohydrate, vitamins, minerals, cofactors and other nutritional components. The present sequence is a genomic DNA encoding a partial novel secreted protein of the invention.

Query Match 58.4%; Score 584.6; DB 22; Length 4977;
Best Local Similarity 74.9%; Pred. No. 4.3e-178;
Matches 745; Conservative 0; Mismatches 249; Indels 1; Gaps 1;

QY 7 GGATAAACCGTGTGAGTGGCCCTCAAGTTGTGTGGCAGCATGGAATGGGAGATGGAGGGA 66
Db 2587 GGCACAGCCATGTGGTGGCCCTCAAGTTGTGTGGCAGCATGGAATGGGAGATGGAGGGA 2646

QY 67 TACATGGATCCCACTACAGGCCAGCTCCCTCAGTATGAGCATGAGCCAGTCAATCT 126
Db 2647 CCAGGGTGGCCCACTAGGGGCCAGTCCCTCTGTAGAGTATGAGTCACTGAGCCT 2706

QY 127 GAATGTGAAGATGGAATGAAGACCGACGAGTGCACATGAGCTCAACCCCTCAATACATG 186
Db 2707 GAGTGCARAAGATGGAGAGGCCAACCCAGAGTCTATGATGACATCAACCCCATACCTG 2766

QY 187 GGGTCAGATCAAGAAACACACACAGAGCTGAGAACTGGTGTAGTGCAGGGTCAGGC 246
Db 2767 GGGACAACCTCAAGAAACACACACAGAGCTGAGAACTGAGTGTGTCAGGACAGGC 2826

QY 247 AAAAACCCCTGACTCCATCTTTATGGCCATGCTAGCTGAATATCCTGTGCAGTATGAT 306
Db 2827 -AAAAACCCCTGATTCATGCTCTTGGCCATGTTAGCCATAATGCTGTGACACTACGTT 2885

QY 307 TTTCTGTGAGAGCAAAACATATTGGCCATATTTTCTTAACCCACCGGTAGTGTGATC 366
Db 2886 TCCCTGTGCACAGCAAAACATATTGGCCATATTTTCCCATCCCCAGCAGTATGGCC 2945

QY 367 ATACTCTGAAGCAGCACTCCTCCTGAGATATATGATCAAGGAGCATCAGTACCAGGA 426
Db 2946 TATACTTTGGAGTCACTCCTCCTGAGATTTATCAGATCAGGGAGAGTGGCTCCAGGA 3005

QY 427 CCTCTAACTCCCTGCACACAGCAATAGACTCTCATACAAATGGTATCAATTATACC 486
Db 3006 CCCCTAACTCCCGTGACATAGAAAGTGTAGACTCTCAGAACAAATGTCAATTAATTATACC 3065

QY 487 ACTCCATTGGAGGACTCTCTTTATGTGTGACCCAGGATACATTTGCTCAACTGTCAGTGC 546
Db 3066 ACTCCACTGGAAGGACTCCCTTTGTTATCACCACAAAGACGCTGCTCAGCCATAGCTGT 3125

QY 547 CTTGAGATTTGATCCCAAGCAGTGGTGTAGTTACCATATAAAAAATATGTACCTATTAGAC 606
Db 3126 CTTGCAATTCAGAGCTCAACATAGTTGTAGTCACTACTATGAAAAAATATGTACTATTAGT 3185

QY 607 CTTAGCTTTATTAATATTACTTGTGTAGTACTAATCACTCTCTGCCCCCATCAACCAAT 666
Db 3186 CTTGGTTCATTAAATGTAAGTGGTGTGCTTAACCAATCATTTCCAGTCCAGTCAACCTAAT 3245

QY 667 TGTACTGATTATACAGATGGGCTCCCTTTGTATTAATTTCTCAACCCCTCTCTTGGGCCAC 726
Db 3246 TGTGCTGATTATACAGATGGATTCATTCATAGTTCTTCCATCCCACTCTGTGGACCCAG 3305

QY 727 TGTCTTGGCCCTTAGCTAGACAATAGTCCATGTTAAATGGGAGACATTAATTGACTGGGT 786

Db 3306 TGTCTTGTATCCACGTGGCTAGTAATAACAATATATGTCAACTGAAGACACTGTGGATTGGGAA 3365
QY 787 CCCTGTGTGTCATTAAAGATGGGAGAGATGAGAATCAGACACATGATGCAATAAACTTCACTGG 846
Db 3366 CTAAGAGTCAATTAGATGGAAAGGTGAAGTCAAGATCATGGCAACAACCTTCACTGG 3425

QY 847 CACTGTGTGGCGAAACCTTTAAACATCTCTTCACTTCAACACACTGGGATTCATCCCAATCT 906
Db 3426 CATTTGGCGCAAGCTTTTAAATGCTTCTTCTTTATACAACAGCAGAAATCCAATCCCAGTCT 3485

QY 907 GCCATGCAACTTGTGTGGCATGGAAACGGCTTTAGCCACACTTTGCCCTCAATGGCATTAAT 966
Db 3486 GCTGCTCAGATGCTTGTGCATGGAGCAGGCTTTAGCCCACTCTTCTCCTCAGTTGCATTAT 3545

QY 967 CAAGGAAGAGAGGTCCTCAATTCAGGAGTCTATGTG 1001
Db 3546 CTGGGAGGAAGGACCAATTCAGAAACTATATG 3580

RESULT 3
AAC79006
ID AAC79006 standard; DNA; 2202 BP.
XX AAC79006;
XX
XX 14-FEB-2001 (first entry)
XX Human secreted protein gene 10 clone HONAH67.
XX
XX Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral;
KW anti-allergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer;
KW vulnery; anticonvulsant; antibacterial; antifungal; antiparasitic;
KW cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
KW neurological disease; infection; human; secreted protein; ss.
XX Homo sapiens.
XX WO200058358-A1.
XX
XX 05-OCT-2000.
XX
XX 23-MAR-2000; 2000WO-US07725.
XX
XX 26-MAR-1999; 99US-0126602.
PR 14-JAN-2000; 2000US-0176063.
XX
XX (HUMA-) HUMAN GENOME SCI INC.
XX
XX Rosen CA, Ruben SM, Komatsoulis G;
PI WPI; 2000-594640/56.
DR P-PSDB; AAB44344.
XX
XX Fourty nine nucleic acid molecules encoding human secreted proteins,
PT useful in the prevention, treatment and diagnosis of cancer, immune
PT disorders, cardiovascular disorders and neurological diseases -
XX
XX Claim 1; Page 321; 367pp; English.
XX
XX The invention relates to the isolation of genes AAC78997-C79045 encoding
CC 49 human secreted proteins AAB44335-B44382. The genes can be used to
CC generate fusion proteins by linking to the gene for the human
CC immunoglobulin G Fc portion (AAC78988) for increasing the stability of
CC the fusion protein as compared to the human protein only. The genes and
CC proteins are useful for preventing, ameliorating or treating medical
CC conditions, e.g. by protein or gene therapy. The genes are isolated
CC from a range of human tissues disclosed in the specification. The
CC nucleic acids, proteins, antibodies and (ant)agonists are useful in
CC the diagnosis, treatment and prevention of: (a) cancer, e.g. breast
CC and ovarian cancer, and other cancers of the adrenal gland, bone, bone
CC marrow, breast, gastrointestinal tract, liver, lung, or urogenital;
CC (b) immune disorders e.g. Addison's disease, allergies, autoimmune

```
CC haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's
CC disease, multiple sclerosis, rheumatoid arthritis and ulcerative
CC colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d)
CC wound healing; (e) neurological diseases e.g. cerebral anoxia and
CC epilepsy; and (f) infectious diseases such as viral, bacterial, fungal
CC and parasitic infections.
XX
SQ Sequence 2202 BP; 633 A; 491 C; 437 G; 633 T; 8 other;

Query Match 45.5%; Score 455.6; DB 21; Length 2202;
Best Local Similarity 76.2%; Pred. No. 1.7e-136;
Matches 571; Conservative 1; Mismatches 176; Indels 1; Gaps 1;

Qy 254 CCTGACTCCATGTTTATGGCCATGCTAGCTGTAATATCTCTGTCAGTATGATTTTTCGT 313
Db 9 CCTGATCCATGTTCTTGGCCATGTTAGCCATATATCTCTGTCAGTATGTTTTCGT 68
Qy 314 GCAGAGCAAAACATATTTGGCCATATTTCTCAACCCAGCGGTAGTGTGA-TCATACTC 372
Db 69 GCAGAGCAAAACATATTTGGCCATATTTCTCAACCCAGCGGTAGTGTGA-TCATACTC 128
Qy 373 TGAAGCAGCACTCTCTGAGATATATCATCATCAAGAGCATCAGTACCAGGACCTCTA 432
Db 129 TGGAGTGACACTCTCTTAAGATTATCATGATTAAAGAGCATGGGCTCCAGGACCCCTA 188
Qy 433 ACTCCCTGTACACAGCAATTTAGACTCTCATACAAATGATCAATTATACCACTCCA 492
Db 189 ACTCCACTGACATAGACAGTGTAGACTCTCAGATAATATGTCATTAATATACCGCTCCA 248
Qy 493 TTGAGGAGCACTCTTTATGTGTCAACCAAGATATGTCACACTGCACTGTCCTTGA 552
Db 249 TTGGAANGACATCTCTTTGTGTGTCAACCAAGATATGTCACACTGCACTGTCCTTGA 308
Qy 553 GTTTGATCCCAAGCATGGTTCAGTTACCATATAAAATATGTCACCTATTAGACCTTAGC 612
Db 309 GTTCAAGCTCACATGGTTCAGTCACTATGGGAAATCATGTACTTATTAAGTCTTGGT 368
Qy 613 TTTATTAATATTACTTGTGTAGTACTAATCACTCTGGCCCACTACCCAAATGTACT 672
Db 369 TATATTAATGTAACGGGTGTCTAACCAACCAATTCCTGGCCCACTGCGCTTCATGTGCT 428
Qy 673 GATTATACAGAAATGGGTCCTTTGATTAATTCACCCCTCTCTGGGCCCCACTGTCTT 732
Db 429 GACTATACAGAAATGGGTCCTTTGATTAATTCACCCCTCTCTGGGCCCCACTGTCTT 488
Qy 733 GGCCCTTAGCTAGACATAGTCTCATGTTAATGGGAGACATTAATGACTGGGGTCCCTGT 792
Db 489 AGCCCACTGGCTAGAAACATCTATGTTAATGGGAGACATTAATGACTGGGGTCCCTGT 548
Qy 793 GGTCATTAAAGTGGAGAGATGAGAATCAGACACATGGCATAAATCTCACTGGCACTGG 852
Db 549 GGCCCAATTAGATGGAAGAAGAAATCAGAAATCATGGCACAATCTTGTCTGGCATGG 608
Qy 853 TGGGCAACTTTTACATCTTCTACTTCAACACACTGGGATTCATATCCCAATCTGCCATG 912
Db 609 TGGCAAGCTTTTAAATGCTTCTCTTTATATAACACTGGGATTCATATCCCAATCTGCCATG 668
Qy 913 CAACCTGCTGGCATGGAACGGCTTTAGCCCACTTTGCTCAATGCGCATTTATCAAGGA 972
Db 669 CAGATGCTTGGCATGGAACGGCTTTAGCCCACTTTGCTCAATGCGCATTTATCAAGGA 728
Qy 973 AAGAGAGTCCCAATTCAGGAGTCTATGTG 1001
Db 729 AGGAAGGACCAATTCAAAAGATGATATG 757

RESULT 4
AAH18610
ID AAH18610 standard; cDNA; 2166 BP.
XX
AC AAH18610;
XX
XX 26-JUN-2001 (first entry)
```

```
XX
DE Human cDNA sequence SEQ ID NO:18816.
XX
KW Human; primer; detection; diagnosis; antisense therapy; gene therapy; ss.
XX
OS Homo sapiens.
XX
FN EPI074617-A2.
XX
PD 07-FEB-2001.
XX
PF 28-JUL-2000; 2000EP-0116126.
XX
PR 29-JUL-1999; 99JP-0248036.
PR 27-AUG-1999; 99JP-0300253.
PR 11-JAN-2000; 2000JP-0118776.
PR 02-MAY-2000; 2000JP-0183767.
PR 09-JUN-2000; 2000JP-0241899.
PA (HELI-) HELIX RES INST.
XX
PI Ota T, Isogai T, Nishikawa T, Hayashi K, Saito K, Yamamoto J;
PI Ishii S, Sugiyama T, Wakamatsu A, Nagai K, Otsuki T;
XX
WI: 2001-318749/34.
XX
PR Primer sets for synthesizing polynucleotides, particularly the 5602
PT full-length cDNAs defined in the specification, and for the detection
PT and/or diagnosis of the abnormality of the proteins encoded by the
PT full-length cDNAs.
XX
PS Claim 8; SEQ ID 18816; 2537pp + CD ROM; English.
XX
CC The present invention describes primer sets for synthesising 5602
CC full-length cDNAs defined in the specification. Where a primer set
CC comprises: (a) an oligo-dT primer and an oligonucleotide complementary
CC to the complementary strand of a polynucleotide which comprises one of
CC the 5602 nucleotide sequences defined in the specification, where the
CC oligonucleotide comprises at least 15 nucleotides; or (b) a combination
CC of an oligonucleotide comprising a sequence complementary to the
CC complementary strand of a polynucleotide which comprises a 5'-end
CC sequence and an oligonucleotide comprising a sequence complementary to a
CC polynucleotide which comprises a 3'-end sequence, where the
CC oligonucleotide comprises at least 15 nucleotides and the combination of
CC the 5'-end sequence/3'-end sequence is selected from those defined in
CC the specification. The primer sets can be used in antisense therapy and
CC in gene therapy. The primers are useful for synthesising polynucleotides,
CC particularly full-length cDNAs. The primers are also useful for the
CC detection and/or diagnosis of the abnormality of the proteins encoded by
CC the full-length cDNAs. The primers allow obtaining of the full-length
CC cDNAs easily without any specialised methods. AAH03166 to AAH13628 and
CC AAH13633 to AAH18742 represent human cDNA sequences; AAH92446 to
CC AAH95893 represent human amino acid sequences; and AAH13629 to AAH13632
CC represent oligonucleotides, all of which are used in the exemplification
CC of the present invention.
XX
SQ Sequence 2166 BP; 611 A; 488 C; 434 G; 633 T; 0 other;

Query Match 45.4%; Score 454; DB 22; Length 2166;
Best Local Similarity 76.4%; Pred. No. 5.4e-136;
Matches 570; Conservative 0; Mismatches 175; Indels 1; Gaps 1;

Qy 257 GACTCCATGTTTATGGCCATGCTAGCTGTAATATCTCTGTCAGTATGATTTTCGTGCA 316
Db 1 GATTCCTGTTCTTGGCCATGTTAGCCATATATCTCTGTCAGTATGTTTTCGTGCA 60
Qy 317 GAAGCAAAACATATTTGGCATATTTTCTCAACCCAGCGGTAGTGTGA-TCATACTCTGA 375
Db 61 GAGCAAAACATATTTGGCATATTTTCTCAACCCAGCGGTAGTGTGA-TCATACTCTGA 120
Qy 376 AGCAGCACTCTCTCTGAGATATATCATCATCAAGAGCATCAGTACCAGGACCTTACT 435
Db 121 AGTGACACTCTCTCTGAGATATATCATCATCAAGAGCATCAGTACCAGGACCTTACT 180
```

QY 436 CCCCTGACACAGACAATTTAGACTCTCTAACAATGATGATCAATTAATATACCACCTCAATG 495
DB 181 CCACCTGACATAGAACAGTTAGACTCTCTCAGAAATATGTCATTAATATACCGCTCAATG 240
QY 496 GAGGACACTTCCTTTATGTCACCCAGAGATACATGCTCAACTGCGATGGCTTGCAGTT 555
DB 241 GAAGGACTTCCTTTGTCACCAACAAGACATCATCAGCCATAGCTGCTTACAGTT 300
QY 556 TGATCCCAAGCATGTTGAGTTACCATAAAAAATATGTACCTATTAGACTTAGCTTT 615
DB 301 CAAGCTCACATAGTTGAGTCATATGCGAAATCATGTACTTATTAAGTCTTGGTTAT 360
QY 616 ATTAATATTACTTGTGTAGTTACTAATACATCTCTGGCCCATCATCCCAAAATTTGACTGAT 675
DB 361 ATTAATGTAACCGGTGTCTTAACCAACCATCTCTGGCCCAATCGCTTCAATTTGCTGAC 420
QY 676 TATACAGAAATGGGCTCCCTTTGATTAATCTCACCCCTCTCTGGCCCATGCTTGGC 735
DB 421 TATACAGAAATGGGCTCCCTTCAATAGTTCTACCCCTCTCCATAGACCATGCTTGGC 480
QY 736 CCTTTAGCTAGACAATAGTCCATGTTAATGGGAGACATTTAGTACTGGGGTCCCTGTGTT 795
DB 481 CACTGGCTAGAAACAATCTATGTTAACTGGAGACATTTGGATTTGGGACCTTAAGGC 540
QY 796 CATTAAAGTGGGAGAGATCAGAATCAGACCATGCGCATAACTTCACTGGCAGCTGGTG 855
DB 541 CAATTAGATGAAAGAAAGAAATCAGAAATCGTGGCACAACCTTTGCTGGCATTGGTG 600
QY 856 CGAACTTTAATCATCTTCTACCTCAACACACTGGGATTCATCCCAATCTGCGCAATGCA 915
DB 601 CAAGCTTTAATGCTTCTTTATATACACTGGGATCCCAATCCAGTCGGCGCCCGAG 660
QY 916 CTTCCTGCGATGGAACGGCTTTAGCCACCTTTGCTCAATGCATTAACAAGAAAG 975
DB 661 ATTCCTGGCATGGAGAGGCTTTAGCCGCTCTCTCTCAGTGGCATTAATAGGGAGG 720
QY 976 AGAGGTCCAATTCAGGAGTCTATG 1001
DB 721 AAGGACCAATTCAAAGATGATG 746
RESULT 5
AAH07625
ID AAH07625 standard; cDNA; 808 BP.
XX AC AAH07625;
XX XX
XX DT 26-JUN-2001 (first entry)
XX XX
XX DE Human cDNA clone (5'-primer) SEQ ID NO:4460.
XX KW Human; primer; detection; diagnosis; antisense therapy; gene therapy; ss.
XX OS Homo sapiens.
XX XX
XX PN EP1074617-A2.
XX XX
XX PD 07-FEB-2001.
XX PF 28-JUL-2000; 2000EP-0116126.
XX XX
XX PR 29-JUL-1999; 99JP-0248036.
XX PR 27-AUG-1999; 99JP-0300253.
XX PR 11-JAN-2000; 2000JP-0118776.
XX PR 02-MAY-2000; 2000JP-0183767.
XX PR 09-JUN-2000; 2000JP-0241899.
XX XX
XX (HELI-) HELIX RES INST.
XX PA
XX PI Ota T, Isogai T, Nishikawa T, Hayashi K, Saito K, Yamamoto J;
XX PI Ishii S, Sugiyama T, Wakamatsu A, Nagai K, Otsuki T;

DR WPI; 2001-318749/34.
XX PT primer sets for synthesizing polynucleotides, particularly the 5602
PT full-length cDNAs defined in the specification, and for the detection
PT and/or diagnosis of the abnormality of the proteins encoded by the
PT full-length cDNAs -
XX PS Claim 1; SEQ ID 4460; 2537pp + CD ROM; English.
XX CC The present invention describes primer sets for synthesizing 5602
CC full-length cDNAs defined in the specification. Where a primer set
CC comprises: (a) an oligo-dT primer and an oligonucleotide complementary
CC to the complementary strand of a polynucleotide which comprises one of
CC the 5602 nucleotide sequences defined in the specification, where the
CC of an oligonucleotide comprising a sequence complementary to the
CC complementary strand of a polynucleotide which comprises a 5'-end
CC sequence and an oligonucleotide comprising a sequence complementary to a
CC polynucleotide which comprises a 3'-end sequence, where the
CC oligonucleotide comprises at least 15 nucleotides and the combination of
CC the 5'-end sequence/3'-end sequence is selected from those defined in
CC the specification. The primer sets can be used in antisense therapy and
CC in gene therapy. The primers are useful for synthesizing polynucleotides,
CC particularly full-length cDNAs. The primers are also useful for the
CC detection and/or diagnosis of the abnormality of the proteins encoded by
CC the full-length cDNAs. The primers allow obtaining of the full-length
CC cDNAs easily without any specialised methods. AAH03166 to AAH13628 and
CC AAH13633 to AAH18742 represent human cDNA sequences; AAB92446 to
CC AAB95993 represent human amino acid sequences; and AAH13629 to AAH13632
CC represent oligonucleotides, all of which are used in the exemplification
CC of the present invention.
XX SQ Sequence 808 BP; 227 A; 186 C; 169 G; 219 T; 7 other;
Query Match 41.8%; Score 418.6; DB 22; Length 808;
Best Local Similarity 76.1%; Pred. No. 9.1e-125;
Matches 553; Conservative 0; Mismatches 171; Indels 3; Gaps 3;
QY 257 GACTCCATGTTTATGGCCATGCTAGCTGTATATATCTCTGTCAGTATGTTTCTGTGCA 316
DB 1 GATTCGCTGTTCTTGGCCATGTTAGCCATATATCTCTGTCAGTATGTTTCTGTGCA 60
QY 317 GAAGCAAAACATATTTGGCATATTTTCTTAACCCACCGGTAGTGTGA-TCATACTCTGA 375
DB 61 GAGGCAAAACATATTTGGCATATTTTCTTAACCCACCGGTAGTGTGA-TCATACTCTGA 120
QY 376 AGCAGCACTCTCTCTGAGATATATCATGATCAAGGAGCATAGTACCAGGACCTTAAT 435
DB 121 AGTGACACTCTCTCTAAGATTTATCATGATTAGGAGCATAGTGGCTCCAGGACCTTAAT 180
QY 436 CCCCTGACACAGACAATTTAGACTCTCTAACAATGATGATCAATTAATACCACCTCAATG 495
DB 181 CCACCTGACATAGAACAGTTAGACTCTCAGAAATATGTCATTAATATACCGCTCAATG 240
QY 496 GAGGACTTCCTTTATGTCACCCAGAGATACATGCTCAACTGCGATGGCTTGCAGTT 555
DB 241 GAAGGACTTCCTTTGTCACCAACAAGACATCATCAGCCATAGCTGCTTACAGTT 300
QY 556 TGATCCCAAGCATGTTGAGTTACCATAAAAAATATGTACCTATTAGACTTAGCTTT 615
DB 301 CAAGCTCACATAGTTGAGTCATATGCGAAATCATGTACTTATTAAGTCTTGGTTAT 360
QY 616 ATTAATATTACTTGTGTAGTTACTAATACATCTCTGGCCCATCATCCCAAAATTTGACTGAT 675
DB 361 ATTAATGTAACCGGTGTCTTAACCAACCATCTCTGGCCCAATCGCTTCAATTTGCTGAC 420
QY 676 TATACAGAAATGGGCTCCCTTTGATTAATCTCACCCCTCTCTGGCCCATGCTTGGC 735
DB 421 TATACAGAAATGGGCTCCCTTCAATAGTTCTACCCCTCTCCATAGACCATGCTTGGC 480
QY 736 CCTTTAGCTAGACAATAGTCCATGTTAATGGGAGACATTTAGTACTGGGGTCCCTGTGTT 795
DB 481 CACTGGCTAGAAACAATCTATGTTAACTGGAGACATTTGGATTTGGGACCTTAAGGC 540

CC	potentially preventing breast cancer. The polynucleotides and encoded
CC	polypeptides are also useful for isolating compounds with cytostatic
CC	activity.
xx	
5Q	Sequence 189 BP; 54 A; 51 C; 34 G; 50 T; 0 other;
	Query Match 18.0%; Score 180.4; DB 22; Length 189;
	Best Local Similarity 99.5%; Pred. No. 5.7e-48;
	Matches 181; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
Qy	418 GTACCAGGACCTTAACCTCCCTTGACACAGAGCAATTAGACTCTCATAAACAATGGTATC 477
Db	3 GTACCAGGACCTTAACCTCCCTTGACACAGAGCAATTAGACTCCCAATGATGTC 62
Qy	478 AATTATACCACTTCATTGGAGGACTTCCTTTATGTGTGCACCGAGATACATTGCTCAAC 537
Db	63 AATTATACCACTTCATTGGAGGACTTCCTTTATGTGTGCACCGAGATACATTGCTCAAC 122
Qy	538 TGCAGTTGCCCTTGCAGTTTGATCCCAAGCATGGTTGAGTTACCATAAAAAAATTATGTAC 597
Db	123 TGCAGTTGCCCTTGCAGTTTGATCCCAAGCATGGTTGAGTTACCAATAAAAAAATTATGTAC 182
Qy	598 CT 599
Db	183 CT 184

RESULT 9	
AAAL18160	standard; cDNA; 189 BP.
XX	
XX	AAAL18160;
XX	
DT	07-DEC-2001 (first entry)
XX	
DE	Human breast cancer expressed polynucleotide 10617.
XX	
KW	Human; breast cancer; cell marker; cytostatic; ss.
XX	
OS	Homo sapiens.
XX	
FN	WO200151628-A2.
XX	
PD	19-JUL-2001.
XX	
PF	10-JAN-2001; 2001WO-US00798.
XX	
PR	14-JAN-2000; 2000US-0176077.
XX	
PR	14-MAR-2000; 2000US-0189167.
XX	
PR	24-MAR-2000; 2000US-0192099.
XX	
PR	29-MAR-2000; 2000US-0193480.
XX	
PR	15-MAY-2000; 2000US-0205230.
XX	
PR	09-JUN-2000; 2000US-0211315.
XX	
PR	25-JUL-2000; 2000US-0220534.
XX	
PA	(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
XX	
PI	Lillie J, Xu Y, Wang Y, Steinmann K;
XX	
DR	WPI; 2001-451856/48.
XX	
PT	New peptide useful as a marker for the diagnosis of breast cancer
XX	
PS	Claim 1; Page 1897; 3695pp; English.
XX	
CC	The invention relates to human breast cancer expressed polynucleotides
CC	(AAAL07544-AAAL26789) and methods of assessing whether a patient is
CC	afflicted with breast cancer by examining the correlation between the
CC	expression of certain markers and the cancerous state of breast cells.
CC	The polynucleotides and encoded polypeptides are potential markers for
CC	detecting, diagnosing, monitoring, characterising treating and
CC	potentially preventing breast cancer. The polynucleotides and encoded
CC	polypeptides are also useful for isolating compounds with cytostatic

CC activity.
XX
SQ Sequence 189 BP; 54 A; 51 C; 34 G; 50 T; 0 other;
Query Match 18.0%; Score 180.4; DB 22; Length 189;
Best Local Similarity 99.9%; Pred. No. 5.7e-48;
Matches 181; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
QY 418 GTACCAGGACCTCTAACTCCCTCGACACAGAGCAATTAGACTCTCTATAACAATGGTATC 477
DB 3 GTACCAGGACCTCTAACTCCCTCGACACAGAGCAATTAGACTCTCTATAACAATGGTATC 62
QY 478 AATTATACCACTCCATTGGAGGACTTCCCTTTATGTCTACCCAGGATACATTGCTCAAC 537
DB 63 AATTATACCACTCCATTGGAGGACTTCCCTTTATGTCTACCCAGGATACATTGCTCAAC 122
QY 538 TGCAGTTGCCCTTCAGTTTCCATCCCAAGCATGGTTGAGTTACCAATAAAAAAATTATGTAC 597
DB 123 TGCAGTTGCCCTTCAGTTTCCATCCCAAGCATGGTTGAGTTACCAATAAAAAAATTATGTAC 182
QY 598 CT 599
DB 183 CT 184
RESULT 10
AAL19385
ID AAL19385 standard; cdna; 184 BP.
AC AAL19385;
XX
XX 07-DEC-2001 (first entry)
XX Human breast cancer expressed polynucleotide 11842.
XX Human; breast cancer; cell marker; cytostatic; ss.
XX Homo sapiens.
XX WO200151628-A2.
XX 19-JUL-2001.
XX 10-JAN-2001; 2001WO-US00798.
XX 14-JAN-2000; 2000US-0176077.
XX 14-MAR-2000; 2000US-0189167.
XX 24-MAR-2000; 2000US-0192099.
XX 29-MAR-2000; 2000US-0193480.
XX 15-MAY-2000; 2000US-0205230.
XX 09-JUN-2000; 2000US-0211315.
XX 25-JUL-2000; 2000US-0220534.
XX (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
PA Lillie J, Xu Y, Wang Y, Steinmann K;
PI WPI; 2001-451856/48.
XX New peptide useful as a marker for the diagnosis of breast cancer -
PT Claim 1; Page 2104; 3695pp; English.
XX The invention relates to human breast cancer expressed polynucleotides
XX (AAL07544-AAL26789) and methods of assessing whether a patient is
XX afflicted with breast cancer by examining the correlation between the
XX expression of certain markers and the cancerous state of breast cells.
XX The polynucleotides and encoded polypeptides are potential markers for
XX detecting, diagnosing, monitoring, characterising treating and
XX potentially preventing breast cancer. The polynucleotides and encoded
XX polypeptides are also useful for isolating compounds with cytostatic
XX activity.

SQ Sequence 184 BP; 54 A; 49 C; 32 G; 49 T; 0 other;
Query Match 17.9%; Score 178.8; DB 22; Length 184;
Best Local Similarity 98.9%; Pred. No. 1.8e-47;
Matches 180; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
QY 418 GTACCAGGACCTCTAACTCCCTCGACACAGAGCAATTAGACTCTCTATAACAATGGTATC 477
DB 3 GTACCAGGACCTCTAACTCCCTCGACACAGAGCAATTAGACTCTCTATAACAATGGTATC 62
QY 478 AATTATACCACTCCATTGGAGGACTTCCCTTTATGTCTACCCAGGATACATTGCTCAAC 537
DB 63 AATTATACCACTCCATTGGAGGACTTCCCTTTATGTCTACCCAGGATACATTGCTCAAC 122
QY 538 TGCAGTTGCCCTTCAGTTTCCATCCCAAGCATGGTTGAGTTACCAATAAAAAAATTATGTAC 597
DB 123 TGCAGTTGCCCTTCAGTTTCCATCCCAAGCATGGTTGAGTTACCAATAAAAAAATTATGTAC 182
QY 598 CT 599
DB 183 CT 184
RESULT 11
AAL26090
ID AAL26090 standard; cdna; 176 BP.
AC AAL26090;
XX
XX 07-DEC-2001 (first entry)
XX Human breast cancer expressed polynucleotide 18547.
XX Human; breast cancer; cell marker; cytostatic; ss.
XX Homo sapiens.
XX WO200151628-A2.
XX 19-JUL-2001.
XX 10-JAN-2001; 2001WO-US00798.
XX 14-JAN-2000; 2000US-0176077.
XX 14-MAR-2000; 2000US-0189167.
XX 24-MAR-2000; 2000US-0192099.
XX 29-MAR-2000; 2000US-0193480.
XX 15-MAY-2000; 2000US-0205230.
XX 09-JUN-2000; 2000US-0211315.
XX 25-JUL-2000; 2000US-0220534.
XX (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
PA Lillie J, Xu Y, Wang Y, Steinmann K;
PI WPI; 2001-451856/48.
XX New peptide useful as a marker for the diagnosis of breast cancer -
PT Claim 1; Page 3419; 3695pp; English.
XX The invention relates to human breast cancer expressed polynucleotides
XX (AAL07544-AAL26789) and methods of assessing whether a patient is
XX afflicted with breast cancer by examining the correlation between the
XX expression of certain markers and the cancerous state of breast cells.
XX The polynucleotides and encoded polypeptides are potential markers for
XX detecting, diagnosing, monitoring, characterising treating and
XX potentially preventing breast cancer. The polynucleotides and encoded
XX polypeptides are also useful for isolating compounds with cytostatic
XX activity.

QY 254 CCGTACTCCATGTTATGGCCATGCTAGCTGTAAATATCCTGTGACGATGATTTTCTGT 313
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 9 CCTGATTCCATGTTCTTGGCCATGTTAGCCATAATATCCTGTGACGATGATTTTCTGT 68
QY 314 GCAGAACCAAAACATATGGGCATATTTTCCCTAACCCACCGGTAGTGCA-TCATACCTC 372
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 69 GCAGAACCAAAACATATGGGCATATGTTNCCCAAGGCCCCAGCAGTATGACCCATACCT 128
QY 373 TGAAGCAGCACTCCTCTCGATATATCATCATCAAGGAGCATCAGTACCA-GGACCTCT 431
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 129 TGGAGTGACACTCTCTCTAGATTTATCATGATTAAGGACATGGGCTCCAGGACCCCT 188
QY 432 AACTCCCTCCGACACAGACAATAGACTCTCTATAACAATGGTATCAATATACCACTCC 491
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 189 AACTCCACCTTGACATAGAACAGTTAGACTCTCAGATAATGTCATTAATATACCGTTCC 248
QY 492 ATTGGAGGGACTTCCTT 508
IIIIII IIIIIII IIIIIII
Db 249 ATTGAAGGACTTCCTT 265
RESULT 14
AAS92505
ID AAS92505 standard; cDNA; 660 BP.
XX
AC AAS92505;
XX
DT 13-FEB-2002 (first entry)
XX
DE DNA encoding novel human diagnostic protein #28309.
XX
KW Human; chromosome mapping; gene mapping; gene therapy; forensic;
KW food supplement; medical imaging; diagnostic; genetic disorder; ss.
XX
OS Homo sapiens.
XX
PN WO200175067-A2.
XX
PD 11-OCT-2001.
XX
PF 30-MAR-2001; 2001WO-US08631.
XX
PR 31-MAR-2000; 2000US-0540217.
XX
PR 23-AUG-2000; 2000US-0649167.
XX
PA (HYSE-) HYSEQ INC.
XX
PI Drmanac RT, Liu C, Tang YT;
XX
DR WPI; 2001-639362/73.
XX
DR P-PSDB; ABG28318.
XX
XX New isolated polynucleotide and encoded polypeptides, useful in
PT diagnostics, forensics, gene mapping, identification of mutations
PT responsible for genetic disorders or other traits and to assess
PT biodiversity -
XX
XX Claim 1; SEQ ID No 28309; 103pp; English.
XX
XX The invention relates to isolated polynucleotide (I) and
XX polypeptide (II) sequences. (I) is useful as hybridisation probes,
CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome
CC and gene mapping, and in recombinant production of (II). The
CC polynucleotides are also used in diagnostics as expressed sequence tags
CC for identifying expressed genes. (I) is useful in gene therapy techniques
CC to restore normal activity of (II) or to treat disease states involving
CC (II). (II) is useful for generating antibodies against it, detecting or
CC quantitating a polypeptide in tissue, as molecular weight markers and as
CC a food supplement. (II) and its binding partners are useful in medical
CC imaging of sites expressing (II). (I) and (II) are useful for treating
CC disorders involving aberrant protein expression or biological activity.
CC The polypeptide and polynucleotide sequences have applications in

CC diagnostics, forensics, gene mapping, identification of mutations
CC responsible for genetic disorders or other traits to assess biodiversity
CC and to produce other types of data and products dependent on DNA and
CC amino acid sequences. AAS64197-AAS94564 represent novel human
CC diagnostic coding sequences of the invention.
CC Note: The sequence data for this patent did not appear in the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 660 BP; 172 A; 168 C; 169 G; 151 T; 0 other;
Query Match 15.3%; Score 152.8; DB 23; Length 660;
Best Local Similarity 82.5%; Pred. No. 1e-38;
Matches 175; Conservative 0; Mismatches 37; Indels 0; Gaps 0;
QY 30 AAGTTGTGTCGACCATGGAATGGAGACTGGAGGATACATGATCCCACTACAGGCC 89
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 353 ATGTTGAGGTGCATCTTGGAAATGGGAGACTGGAGGGACCCATGATCCCACTGACCC 412
QY 90 CAGCTCCTCCAGTATGAGCCATGAGCCAGTTCGAATCTGAATGTGAAGTGAATGAAGAC 149
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 413 TGGTTCCTCCCATCATCATCATGATGATGATGATGATGATGATGATGATGATGATGAT 472
QY 150 CGACGAGAGTCACATGACCTCAACCTCATACATGGGTGATCAAGAAACCAACACAC 209
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 473 TGACGAGAGTCATGCTGACATCAACCCCATACATGGGACTGATCAAGAAACCAACACAC 532
QY 210 CAGAAGCTGAGAACTGGTGTAGTGCAGGGT 241
IIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII IIIIIII
Db 533 AGGAAGCTGAGAACTGCTGGAGTGCAGGGT 564
RESULT 15
AAHL5033
ID AAHL5033 standard; cDNA; 2652 BP.
XX
AC AAHL5033;
XX
DT 26-JUN-2001 (first entry)
XX
DE Human cDNA sequence SEQ ID NO:13009.
XX
KW Human; primer; detection; diagnosis; antisense therapy; gene therapy; ss.
XX
OS Homo sapiens.
XX
PN EP1074617-A2.
XX
PD 07-FEB-2001.
XX
PF 28-JUL-2000; 2000EP-0116126.
XX
PR 29-JUL-1999; 99JP-0248036.
XX
PR 27-AUG-1999; 99JP-0300253.
XX
PR 11-JAN-2000; 2000JP-0118776.
XX
PR 02-MAY-2000; 2000JP-0183767.
XX
PR 09-JUN-2000; 2000JP-0241899.
XX
XX (HELI-) HELIX RES INST.
XX
XX Ota T, Isoqai T, Nishikawa T, Hayashi K, Saito K, Yamamoto J;
PI Ishii S, Sugiyama T, Wakamatsu A, Nagai K, Otsuki T;
XX
XX WPI; 2001-318749/34.
XX
XX Primer sets for synthesizing polynucleotides, particularly the 5602
PT full-length cDNAs defined in the specification, and for the detection
PT and/or diagnosis of the abnormality of the proteins encoded by the
PT full-length cDNAs -
XX
XX Claim 8; SEQ ID 13009; 2537pp + CD ROM; English.
XX
XX The present invention describes primer sets for synthesising 5602

full-length cDNAs defined in the specification. Where a primer set comprises: (a) an oligo-dr primer and an oligonucleotide complementary to the complementary strand of a polynucleotide which comprises one of the 5602 nucleotide sequences defined in the specification, where the oligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprising a sequence complementary to the complementary strand of a polynucleotide which comprises a 5'-end sequence and an oligonucleotide comprising a sequence complementary to a polynucleotide which comprises a 3'-end sequence, where the oligonucleotide comprises at least 15 nucleotides and the combination of the 5'-end sequence/3'-end sequence is selected from those defined in the specification. The primer sets can be used in antisense therapy and in gene therapy. The primers are useful for synthesising polynucleotides, particularly full-length cDNAs. The primers are also useful for the detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs. The primers allow obtaining of the full-length cDNAs easily without any specialised methods. AAH03166 to AAH13628 and AAH13633 to AAH18742 represent human cDNA sequences; AAB92446 to AAB95893 represent human amino acid sequences; and AAH13629 to AAH13632 represent oligonucleotides, all of which are used in the exemplification of the present invention.

XX
SQ Sequence 2652 BP; 818 A; 567 C; 600 G; 667 T; 0 other;

Query Match 15.2%; Score 152.6; DB 22; Length 2652;
Best Local Similarity 68.1%; Pred. No. 2.5e-38;
Matches 228; Conservative 0; Mismatches 104; Indels 3; Gaps 1;
Qy 1 GGAGATGGATAAACCGTGTGAGTGCCTCAAGTTGTGTGCGACCATGGATGGGAGACTG 60
Db 1950 GGAGATGGACAAGCCGTGTGGTGCCTCAAGGTGTGTGCAACCATGGATGGGAGACTG 2009
Qy 61 GAGGATACATGGATCCCACTACAGGCCAGCTCCTCCAGTATGAGCCATGAGCCAGTT 120
Db 2010 GAGGAACCCAGGTGGCCAAACCATGGTCCGGTCCCTCTGTGTGAGCCATGAGCCAGCT 2069
Qy 121 GAATCTGAATGTGAATGGAATGAAGACCCAGCAGAGTCACTGACGTCAACCCCTCAT 180
Db 2070 GAGCCTGAGTCCGAGAGCGGAGAGAGGCCGACCCACAGTCA--TGACATCAACCCCAT 2126
Qy 181 AACATGGGGTCAGATCAAGAAACACACACAGAGCTGAGAACTGGTGTAGTGCACAGG 240
Db 2127 AACCTGGGGACAACCTCAAGAAACACACACAGAGGCTGAGAACTACTGGAGCACCAGG 2186
Qy 241 TCAGCAAAACCCCTGACTCCATGTTTATGGCCATGCTAGCTGTATATCTCTGTGCAGT 300
Db 2187 ACAGTCTGTAAGTTGGATGGACCATCAATGGGAAATGAGAGCTGCCACCCTGGCCCTT 2246
Qy 301 ATGATTTTCTGTGCAGAGCAAAACATATTTGG 335
Db 2247 ACACCTCTTCAATTAAATACATAAACAAGAGGAGG 2281

Search completed: December 24, 2002, 17:17:51
Job time : 181.952 secs

GenCore version 5.1.3
Copyright (c) 1993 - 2002 CompuGen Ltd.

OM nucleic - nucleic search, using sw model

Run on: December 24, 2002, 16:23:56 ; Search time 165.952 Seconds
(without alignments)
13593.723 Million cell updates/sec

Title: US-09-708-724A-3_COPY_99000_100000
Perfect score: 1001
Sequence: 1 tggcagcgctgtagtc...ttttatccaccatcaactaa 1001

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 2185239 seqs, 1125999159 residues
Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : N_Geneseq_101002:*
1: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1980.DAT:*
2: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1981.DAT:*
3: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1982.DAT:*
4: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1983.DAT:*
5: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1984.DAT:*
6: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1985.DAT:*
7: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1986.DAT:*
8: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1987.DAT:*
9: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1988.DAT:*
10: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1989.DAT:*
11: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1990.DAT:*
12: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1991.DAT:*
13: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1992.DAT:*
14: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1993.DAT:*
15: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1994.DAT:*
16: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1995.DAT:*
17: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1996.DAT:*
18: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1997.DAT:*
19: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1998.DAT:*
20: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1999.DAT:*
21: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2000.DAT:*
22: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001A.DAT:*
23: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001B.DAT:*
24: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2002.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Score	Length	ID	Description
1	425.2	42.5	2750	21 AAC69110	Human secreted pro
2	424	42.4	2752	21 AAC69119	Human secreted pro
c 3	314	31.4	2391	22 AA192576	Human polynucleoti
c 4	121.2	12.1	2266	22 AAK89548	Human digestive sy
c 5	119	11.9	16161	22 AAK83469	Human immune/haema
c 6	118.4	11.8	1031	22 AAK82137	Human immune/haema
c 7	117.4	11.7	17792	22 AAS32727	Human genomic DNA
c 8	117.4	11.7	17792	22 AAS36099	Human cardiovascular
c 9	117	11.7	87350	18 AAK83003	Human WRN genomic

c 10	116.8	11.7	1039	22 AAK62290	Human immune/haema
c 11	116.8	11.7	16106	22 AAK83468	Human immune/haema
c 12	115.4	11.5	666	22 AAK64559	Human immune/haema
c 13	114.6	11.4	929	21 AAC59766	Human secreted pro
c 14	114.2	11.4	2361	22 ABA21235	Human nervous syst
c 15	114	11.4	30620	22 AAK66931	Human immune/haema
c 16	114	11.4	325791	22 AAS43104	Human Oestrogen re
c 17	113.8	11.4	734	22 ABA16869	Human nervous syst
c 18	113.8	11.4	734	22 ABA16871	Human nervous syst
c 19	113.8	11.4	734	22 ABA20112	Human nervous syst
c 20	113.8	11.4	734	22 ABA20114	Human nervous syst
c 21	113.8	11.4	1184	21 AAS26411	Human secreted pro
c 22	113.8	11.4	4185	22 AAS21277	Human cDNA sequenc
c 23	113.8	11.4	11987	22 AAL07284	Human reproductiv
c 24	113.8	11.4	11987	23 ABL98830	Human testicular a
c 25	113.8	11.4	46366	22 AAK82098	Human immune/haema
c 26	113.8	11.4	227968	24 AAK83497	Human cDNA differe
c 27	113.4	11.3	6216	24 AKN83166	Human PER2 S62GG
c 28	113.4	11.3	6218	20 AAX58987	Human transcriptio
c 29	113.4	11.3	6219	24 AKN83167	Human PER2 cDNA.
c 30	113.4	11.3	9039	22 AAK75933	Human immune/haema
c 31	113.4	11.3	9039	22 AAK85246	Human immune/haema
c 32	113.4	11.3	35414	21 AAD00147	TR12 related DNA-1
c 33	112.6	11.2	590	24 AKN63415	Human cancer relat
c 34	112.6	11.2	20835	22 AAK86765	Human immune/haema
c 35	112.6	11.2	25806	22 AAK86766	Human immune/haema
c 36	112.4	11.2	2147	23 AKA42851	Genomic sequence #
c 37	112.2	11.2	6708	22 ABA07966	Human ovarian and
c 38	112.2	11.2	6708	22 AAL03878	Human reproductiv
c 39	112.2	11.2	32082	22 AAL06991	Human reproductiv
c 40	112.2	11.2	32186	22 ABA21319	Human nervous syst
c 41	112.2	11.2	32186	22 AAK89692	Human digestive sy
c 42	112.2	11.2	99014	24 AKN96931	Gene #3429 used to
c 43	112	11.2	13493	22 AAK74889	Human immune/haema
c 44	112	11.2	13493	22 AAK82826	Human immune/haema
c 45	112	11.2	35973	24 ABA13076	Human amyloid beta

ALIGNMENTS

RESULT 1
AAC69110
ID AAC69110 standard; DNA; 2750 BP.
XX AC AAC69110;
XX DT 31-JAN-2001 (first entry)
XX Human secreted protein gene 27 clone HOUHD63.
XX Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral;
KW antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer;
KW vulnary; anticonvulsant; antibacterial; antifungal; antiparasitic;
KW cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
KW neurological disease; infection; human; secreted protein; ss.
OS Homo sapiens.
XX WO20005371-A1.
PN 21-SEP-2000.
XX 16-MAR-2000; 2000WO-US06783.
XX 18-MAR-1999; 99US-0125055.
XX (HUMA-) HUMAN GENOME SCI INC.
XX Ruben SM, Ni J, Ebner R, Rosen CA, Shi Y, Birse C, Florence K;
PI Komatsoulis G, Lafleur DW, Moore PA, Olsen HS, Young PE;
XX WPI; 2000-594448/56.
DR

DR	P-PSDB; AAB38010.	
XX	New nucleic acid molecules encoding 27 human secreted proteins for	
PT	diagnosing, preventing, treating or ameliorating medical conditions and	
PT	used as food additives or preservatives -	
XX	Claim 1; Page 382-383; 453pp; English.	
PS	The invention relate to the isolation of genes AAC69084-C69119 encoding	
CC	27 human secreted proteins AAB37984-B38019. The genes can be used to	
CC	generate fusion proteins by linking to the gene for the human	
CC	immunoglobulin G Fc portion (AAC69075) for increasing the stability of	
CC	the fusion protein as compared to the human protein only. The genes and	
CC	proteins are useful for preventing, ameliorating or treating medical	
CC	conditions, e.g. by protein or gene therapy. The genes are isolated	
CC	from a range of human tissues disclosed in the specification. The	
CC	nucleic acids, proteins, antibodies and (ant)agonists are useful in	
CC	the diagnosis, treatment and prevention of: (a) cancer, e.g. breast	
CC	and ovarian cancer, and other cancers of the adrenal gland, bone, bone	
CC	marrow, breast, gastrointestinal tract, liver, lung, or urogenital;	
CC	(b) immune disorders e.g. Addison's disease, allergies, autoimmune	
CC	haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's	
CC	disease, multiple sclerosis, rheumatoid arthritis and ulcerative	
CC	colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d)	
CC	wound healing; (e) neurological diseases e.g. cerebral anoxia and	
CC	epilepsy; and (f) infectious diseases such as viral, bacterial, fungal	
CC	and parasitic infections.	
XX	Sequence 2750 BP; 803 A; 526 C; 593 G; 816 T; 12 other;	
Qy	Query Match 42.58; Score 425.2; DB 21; Length 2750;	
Db	Best Local Similarity 79.08; Pred. No. 6.1e-109;	
XX	Matches 591; Conservative 1; Mismatches 124; Indels 32; Gaps 6;	
Qy	284 GGTGTTGGCGTGGCGGTGTTGCTCTGCTGCACAGGTGGAGTGGAGTGCCTC 343	
Db	158 GCTCTGCGGTGGCGCGGTGCTGCTCTGCTGCGGAGTCGGAATCGGAACGCCCTC 217	
Qy	344 TTCTCTCTCAGACAGAACCATG-AGCCTAGCGGCGAGCGCGGTTCGCGAAGTCCCC 402	
Db	218 GTCTGTCTCAGGCCAGAACCATGAACCGCGGCGGCGGCGCGCGCATGCTCCCT 277	
Qy	403 CTCGCCAACGGCGGCTTCCTCAGAGCGGTCC--GTGCCCCGCTGCGGGAGTGACCC 460	
Db	278 CTCGGCCAGGTGGCGCGCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 337	
Qy	461 CGAGCGAGTCCAGATGTCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 520	
Db	338 CGAGCGAGTCCAGATGTCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 397	
Qy	521 CATCTCCAGGAGCGGCTGAGTAGGAACTGCAGCCGC-----CACATCCTCTC 569	
Db	398 CCTCCCGGGACCCAGCGAGGAGGAGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCTA 457	
Qy	570 TTTACCGGGATGTCAGGATTTACCGTGAATCATGACTCGTCACTCGTCAATTTACCA 629	
Db	458 TTTCTCTGGGATGTCAGGATTTACCATGAATTTATGACTCGTCACTCGTCAATTTATCA 517	
Qy	630 ATGGGAAATTTGGAGTCTAGAAATTTATGCTCTCCATTTTACCCACCGGTTCCCGAGTAG 689	
Db	518 ATGGGAAATTTGGAGTCTAGAAATTTATGCTCTCCATTTTACCCACCGGTTCCCGAGTAG 577	
Qy	690 CTGTATTTGGGTGATGAAGTCTCCGGAA-----CAAATGCGCTGCCATGATAG 738	
Db	578 TTATATTTGGGTGATGAAGTCTCCGGAA-----CAAATGCGCTGCCATGATAG 637	
Qy	739 TTTTCTGAAAGTACATGTTTGGTTTCCAGACACAAATACAGACTCTGGAGCTTTAA 798	
Db	638 TTTTCTGAAAGTACATGTTTGGTTTCCAGACACAAATACAGACTCTGGAGCTTTAA 697	
Qy	799 GCACCTTTATATGTTATTTAGTTAATGCT-----TTTAAAGTCAGAGTACTTTTCAAGGA 853	
Db	698 GCACCTTTATATGTTATTTAGTTAATGCTTTTAAATTTAGTCAGAGTACTTTTCAAGGA 757	

Qy	854 AAATTGGAATGATTGGAATAGGACTCCACAGCATCTAATTGTAGATGTCAATCTTCT 913
Db	758 AAGTTTGAATGTTTGAATAGGACTCCATGACATCACTGTAGA--TCCAGTCCCTCT 815
Qy	914 CATACTACAATCATTTCCAGGAAGGAAAGATAGGACCTTTGAAATAATCTGTGGATCG 973
Db	816 CATACTACAATCATTTCCAGGAAGGAAAGATAGGACCTTTGAAATAATCTGTAGTCT 875
Qy	974 GCCATGTGTTTATCCACCATCACTAA 1001
Db	876 GCCATGTGTTTATCCACCATCACTAA 903
XX	RESULT 2
XX	AAC69119
XX	ID AAC69119 standard; DNA; 2752 BP.
XX	AC AAC69119;
XX	XX AAC69119;
DT	31-JAN-2001 (first entry)
XX	Human secreted protein gene 27 clone HPJBF63.
XX	Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral;
KW	antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer;
KW	vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic;
KW	cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
KW	neurological disease; infection; human; secreted protein; ss.
XX	Homo sapiens.
OS	WO200055371-Al.
XX	21-SEP-2000.
PD	16-MAR-2000; 2000WO-US06783.
PF	18-MAR-1999; 99US-0125055.
XX	(HUMA-) HUMAN GENOME SCI INC.
PA	Ruben SM, Ni J, Ebner R, Rosen CA, Shi Y, Birse C, Florence K;
XX	Komatsoulis G, Lafleur DW, Moore PA, Olsen HS, Young PE;
PI	WPI: 2000-594448/56.
PI	P-PSDB; AAB38019.
XX	New nucleic acid molecules encoding 27 human secreted proteins for
PT	diagnosing, preventing, treating or ameliorating medical conditions and
PT	used as food additives or preservatives -
XX	Claim 1; Page 390-391; 453pp; English.
PS	The invention relate to the isolation of genes AAC69084-C69119 encoding
XX	27 human secreted proteins AAB37984-B38019. The genes can be used to
CC	generate fusion proteins by linking to the gene for the human
CC	immunoglobulin G Fc portion (AAC69075) for increasing the stability of
CC	the fusion protein as compared to the human protein only. The genes and
CC	proteins are useful for preventing, ameliorating or treating medical
CC	conditions, e.g. by protein or gene therapy. The genes are isolated
CC	from a range of human tissues disclosed in the specification. The
CC	nucleic acids, proteins, antibodies and (ant)agonists are useful in
CC	the diagnosis, treatment and prevention of: (a) cancer, e.g. breast
CC	and ovarian cancer, and other cancers of the adrenal gland, bone, bone
CC	marrow, breast, gastrointestinal tract, liver, lung, or urogenital;
CC	(b) immune disorders e.g. Addison's disease, allergies, autoimmune
CC	haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's
CC	disease, multiple sclerosis, rheumatoid arthritis and ulcerative
CC	colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d)
CC	wound healing; (e) neurological diseases e.g. cerebral anoxia and
CC	epilepsy; and (f) infectious diseases such as viral, bacterial, fungal
CC	and parasitic infections.

Query Match 42.4%; Score 424; DB 21; Length 2752;
 Best Local Similarity 79.0%; Pred. No. 1.3e-108;
 Matches 591; Conservative 0; Mismatches 125; Indels 32; Gaps 6;

Qy	284	GGTTTGGGTCGGCGGTGTTGTCCTGCTGCCACAGGTGGAACTGGAGATGCCCTC	343
Db	165	GCTCTGCGGTCGGCGGTGTCCTGCTGCGCAGCTCGGAATCGGAACGCCCTC	224
Qy	344	TTCCCTCTCTCAGACAGAACCATG-AGCCTAGCGGACGCCGGTTCGGAAGCTCCCC	402
Db	225	GTCTGTCTCTCAGGCCAGAACCATGAACCCGGGCGGACGCGCGCGCATGCTCCCT	284
Qy	403	CTCCGCCAACCGGCGCTCTCTCAGAGCGCTCC--GTGCCCGCCCTGCCGGGAGTGCACC	460
Db	285	CTCGCCGAGTGCAGCGCGCTCAGTGCTTCACCGTTCCTGAGCGGAGCC	344
Qy	461	GCAGCGCAGTGCAGAGTTGTCCTGTTGGCGGTGACCAAGGAGGACTGGAGCGCGGAT	520
Db	345	GCAGCGCAGCAACGAATGCTCTGTTGGCGGCGCGGAGGAGTGGAGCGGACGGA	404
Qy	521	CATCTCCAGGAAGCGGCTGAGTAGGAACTCCAGCCG-----CACATCTCTC	569
Db	405	CCTCCCGGACCGACGCAAGAGGAGGACCGACCGCGCGCCGACGATCAGCTCTCA	464
Qy	570	TTTACCGCGGATGTGCAGGATTACCGGTGAATCATGATCGTCTCTCGGAATTACCA	629
Db	465	TTTCCCTGGGATGTGCAGAAATACATGAATATGACTCGTATCTGAGAATATCA	524
Qy	630	ATGGGAAATTTGGAGTCTAGAAATATTTCCTCCATTTAGCCCAACCGGTTCCCGAGTAG	689
Db	525	ATGGGAAACTGGAGTCTAGAAATGTGCTACCATTTTAGCCCAACCGGTTCCCGAGTAG	584
Qy	690	CTGTATTGGGTGATGAAGTCTCCGNA-----CAATGCGCTGCCATGATAG	738
Db	585	TTATATTGGGTGATAAATGTTCCGAAATGCAATTTGCACAAATTCAGTCTGATGACAA	644
Qy	739	TTTCTGAAAAAGTAACATGTTTGGTTTCCAGAACACAAATACAGACTCTGGAGCTTTTAA	798
Db	645	TTTTGTGAAAGTAACACGTTTGGTGGCCCAAGACACAACTACTGACTTTGGAGCTTTTAA	704
Qy	799	GCACCTTTATATGTTATTAGTTAATGCT-----TTTAACTCAGAGTAGTTTATCAAAAGGA	853
Db	705	GCACCTTTATATGTTATTAGTTAATGCTTTTAAATTAAGTCAGAATAGTTTATCAAAAGGA	764
Qy	854	AAATTTGAATGTTGGAATAGGACTCCAGACATCTAATTTGATAGATGCCAATTCCTCT	913
Db	765	AAGTTTGAATGTTTGGAAATAGGACTCCATGATCTAAGTGTAGA--TCCAGTCTCTCT	822
Qy	914	CATACTCAAAATCATTTCCAGGAAGGAAAAAGATAGGACCTTTGAAAAATCTGATGGATCG	973
Db	823	CATCTAGCAATGTTGCCAGGAGAAATGAGGACCTGTGAAAAATCTGATGAGTCT	882
Qy	974	GCCATGTGTTTTTATCCACCATCACTAA	1001
Db	883	GCCATGAGTTTTTATCCACCATCACTAA	910

RESULT 3
 AAI92576/c
 ID AAI92576 standard; cDNA; 2391 BP.
 XX AC AAI92576;
 XX AC AAI92576;
 XX 06-NOV-2001 (first entry)
 XX Human polynucleotide SEQ ID NO 12636.
 XX Human; cytokine; cell proliferation; cell differentiation; gene therapy;
 KW vaccine; peptide therapy; stem cell growth factor; haematopoiesis;
 KW tissue growth factor; immunomodulatory; cancer; leukemia;

KW	nervous system disorders; arthritis; inflammation; ss.
XX	Homo sapiens.
XX	WO200164835-A2.
XX	07-SEP-2001.
PD	
XX	26-FEB-2001; 2001WO-US04927.
XX	
XX	28-FEB-2000; 2000US-0515126.
PR	18-MAY-2000; 2000US-0577409.
XX	(HYSE-) HYSEQ INC.
XX	
XX	Tang YT, Liu C, Drmanac RT;
PI	WPI; 2001-514838/56.
DR	P-PSDB; AAO12645.
XX	
PT	Isolated nucleic acids and polypeptides, useful for preventing
PT	diagnosing and treating e.g. leukaemia, inflammation and immune
PT	disorders -
XX	
PS	Claim 1; SEQ ID NO 12636; 1399pp + Sequence Listing; English.
XX	
CC	The invention relates to human polynucleotides (AAI79941-AAI93841) and
CC	the encoded proteins (AAO0010-AAO13910) that exhibit activity relating to
CC	cytokine, cell proliferation or cell differentiation or which may induce
CC	production of other cytokines in other cell populations. The
CC	polynucleotides and polypeptides are useful in gene therapy, vaccines or
CC	peptide therapy. The polypeptides have various cytokine-like activities,
CC	e.g. stem cell growth factor activity, haematopoiesis regulating
CC	activity, tissue growth factor activity, immunomodulatory activity and
CC	activin/inhibin activity and may be useful in the diagnosis and/or
CC	treatment of cancer, leukaemia, nervous system disorders, arthritis and
CC	inflammation.
CC	Note: The sequence data for this patent did not form part of the printed
CC	specification, but was obtained in electronic format directly from WIPO
CC	at ftp.wipo.int/pub/published_pct_sequences.
XX	
XX	Sequence 2391 BP; 766 A; 480 C; 379 G; 762 T; 4 other;
QY	
Query Match	31.4%; Score 314; DB 22; Length 2391;
Best Local Similarity	81.1%; Pred. No. 1e-77;
Matches	425; Conservative 0; Mismatches 70; Indels 29; Gaps 4;
QY	505 GACTGGAGCGGGATCATCTCCAGGAAGCGGGCTGATAGAACTGACGGCG----- 558
Db	
QY	2359 GACTGGAGCGGACGAGCCTCCCGGGGACCCAGCGAAGGAGGAGCGCGCTTCC 2300
Db	
QY	559 -----CACATCCTCTCTTTACCGGGGATGTCAGGATTACCGTCAATCATGACTCGTC 613
Db	
QY	2299 TTCAATCAGCTCTCTATTTCTCCGGGATGTCAGAAATACCATGAAATATGACTCGTC 2240
Db	
QY	614 ATCTCGCAATTACCAATGGGAAATTCGAGTCTAGAAATTTATTCCTCCATTTAGGCC 673
Db	
QY	2239 ATCTGAGAAATATCAATGGGAACTGGAGTCTAGAAATGTTCTACCATTTAGGCC 2180
Db	
QY	674 ACCGGTTCCTCCAGTAGCTGTATTTGGGGTGATGAAGTGTCTCGGAA-----CAA 722
Db	
QY	2179 ACCGGTTCCTCCCAATAGTTATATTTGGGTGATAAAATGTTCCGAAATGATGTCACAAAT 2120
Db	
QY	723 TGGCGTCCAGTAGATAGTTTCTGAAAGCTTAACATGTTGGTTTCCAGAACACATACAG 782
Db	
QY	2119 TCAGCTGCTATGACAAATTTTGGAAAGTACATCTTTGGTGCCCCAGAACACATCTG 2060
Db	
QY	783 ACTCTGGAGCTTTAAGCACCTTTATATGTTATTTAGTTAAATGCT-----TTTAAGTCAGA 837
Db	
QY	2059 ACTTTGGAGCTTTAAGCACCTTTATATGTTATTTAGTTAAATGCTTTTAAATGTCAGA 2000
Db	
QY	838 GTAGTTTATCAAGGAAATTTGAATGATTTGAATTAAGGACTCCACAGCATCTAATGTA 897
Db	

PR 04-FEB-2000; 2000US-0180628.
PR 24-FEB-2000; 2000US-0184664.
PR 02-MAR-2000; 2000US-0186350.
PR 16-MAR-2000; 2000US-0189874.
PR 17-MAR-2000; 2000US-0190076.
PR 18-APR-2000; 2000US-0198123.
PR 19-MAY-2000; 2000US-0205515.
PR 07-JUN-2000; 2000US-0209467.
PR 30-JUN-2000; 2000US-0214886.
PR 28-JUN-2000; 2000US-0215135.
PR 07-JUL-2000; 2000US-0216647.
PR 07-JUL-2000; 2000US-0216880.
PR 11-JUL-2000; 2000US-0217487.
PR 11-JUL-2000; 2000US-0217496.
PR 14-JUL-2000; 2000US-0218290.
PR 26-JUL-2000; 2000US-0220964.
PR 26-JUL-2000; 2000US-0220964.
PR 14-AUG-2000; 2000US-0224518.
PR 14-AUG-2000; 2000US-0224519.
PR 14-AUG-2000; 2000US-0225213.
PR 14-AUG-2000; 2000US-0225214.
PR 14-AUG-2000; 2000US-0225266.
PR 14-AUG-2000; 2000US-0225267.
PR 14-AUG-2000; 2000US-0225268.
PR 14-AUG-2000; 2000US-0225270.
PR 14-AUG-2000; 2000US-0225447.
PR 14-AUG-2000; 2000US-0225757.
PR 14-AUG-2000; 2000US-0225758.
PR 14-AUG-2000; 2000US-0225759.
PR 18-AUG-2000; 2000US-0226279.
PR 22-AUG-2000; 2000US-0226681.
PR 22-AUG-2000; 2000US-0226688.
PR 22-AUG-2000; 2000US-0227182.
PR 23-AUG-2000; 2000US-0227009.
PR 30-AUG-2000; 2000US-0228924.
PR 01-SEP-2000; 2000US-0229287.
PR 01-SEP-2000; 2000US-0229343.
PR 01-SEP-2000; 2000US-0229344.
PR 01-SEP-2000; 2000US-0229345.
PR 05-SEP-2000; 2000US-0229509.
PR 05-SEP-2000; 2000US-0229513.
PR 06-SEP-2000; 2000US-0230437.
PR 06-SEP-2000; 2000US-0230438.
PR 08-SEP-2000; 2000US-0231242.
PR 08-SEP-2000; 2000US-0231243.
PR 08-SEP-2000; 2000US-0231244.
PR 08-SEP-2000; 2000US-0231244.
PR 08-SEP-2000; 2000US-0231413.
PR 08-SEP-2000; 2000US-0231414.
PR 08-SEP-2000; 2000US-0232080.
PR 08-SEP-2000; 2000US-0232081.
PR 12-SEP-2000; 2000US-0231968.
PR 14-SEP-2000; 2000US-0232397.
PR 14-SEP-2000; 2000US-0232398.
PR 14-SEP-2000; 2000US-0232399.
PR 14-SEP-2000; 2000US-0232401.
PR 14-SEP-2000; 2000US-0232401.
PR 14-SEP-2000; 2000US-0233063.
PR 14-SEP-2000; 2000US-0233064.
PR 14-SEP-2000; 2000US-0233064.
PR 14-SEP-2000; 2000US-0233065.
PR 21-SEP-2000; 2000US-0234223.
PR 21-SEP-2000; 2000US-0234274.
PR 25-SEP-2000; 2000US-0234997.
PR 25-SEP-2000; 2000US-0234998.
PR 26-SEP-2000; 2000US-0235484.
PR 27-SEP-2000; 2000US-0235834.
PR 27-SEP-2000; 2000US-0235836.
PR 29-SEP-2000; 2000US-0236327.
PR 29-SEP-2000; 2000US-0236367.
PR 29-SEP-2000; 2000US-0236368.
PR 29-SEP-2000; 2000US-0236369.
PR 29-SEP-2000; 2000US-0236370.
PR 02-OCT-2000; 2000US-0236802.
PR 02-OCT-2000; 2000US-0237037.

PR 02-OCT-2000; 2000US-0237038.
PR 02-OCT-2000; 2000US-0237039.
PR 02-OCT-2000; 2000US-0237040.
PR 13-OCT-2000; 2000US-0239935.
PR 13-OCT-2000; 2000US-0239937.
PR 20-OCT-2000; 2000US-0240960.
PR 20-OCT-2000; 2000US-0241221.
PR 20-OCT-2000; 2000US-0241785.
PR 20-OCT-2000; 2000US-0241786.
PR 20-OCT-2000; 2000US-0241787.
PR 20-OCT-2000; 2000US-0241808.
PR 20-OCT-2000; 2000US-0241809.
PR 20-OCT-2000; 2000US-0241826.
PR 01-NOV-2000; 2000US-0244617.
PR 08-NOV-2000; 2000US-0246474.
PR 08-NOV-2000; 2000US-0246475.
PR 08-NOV-2000; 2000US-0246476.
PR 08-NOV-2000; 2000US-0246477.
PR 08-NOV-2000; 2000US-0246478.
PR 08-NOV-2000; 2000US-0246523.
PR 08-NOV-2000; 2000US-0246524.
PR 08-NOV-2000; 2000US-0246525.
PR 08-NOV-2000; 2000US-0246526.
PR 08-NOV-2000; 2000US-0246527.
PR 08-NOV-2000; 2000US-0246528.
PR 08-NOV-2000; 2000US-0246532.
PR 08-NOV-2000; 2000US-0246609.
PR 08-NOV-2000; 2000US-0246610.
PR 08-NOV-2000; 2000US-0246611.
PR 08-NOV-2000; 2000US-0246613.
PR 17-NOV-2000; 2000US-0249207.
PR 17-NOV-2000; 2000US-0249208.
PR 17-NOV-2000; 2000US-0249209.
PR 17-NOV-2000; 2000US-0249210.
PR 17-NOV-2000; 2000US-0249211.
PR 17-NOV-2000; 2000US-0249212.
PR 17-NOV-2000; 2000US-0249213.
PR 17-NOV-2000; 2000US-0249214.
PR 17-NOV-2000; 2000US-0249215.
PR 17-NOV-2000; 2000US-0249216.
PR 17-NOV-2000; 2000US-0249217.
PR 17-NOV-2000; 2000US-0249218.
PR 17-NOV-2000; 2000US-0249244.
PR 17-NOV-2000; 2000US-0249245.
PR 17-NOV-2000; 2000US-0249264.
PR 17-NOV-2000; 2000US-0249265.
PR 17-NOV-2000; 2000US-0249297.
PR 17-NOV-2000; 2000US-0249299.
PR 17-NOV-2000; 2000US-0249300.
PR 01-DEC-2000; 2000US-0250160.
PR 01-DEC-2000; 2000US-0250391.
PR 05-DEC-2000; 2000US-0251030.
PR 05-DEC-2000; 2000US-0251988.
PR 05-DEC-2000; 2000US-0256719.
PR 06-DEC-2000; 2000US-0251479.
PR 08-DEC-2000; 2000US-0251856.
PR 08-DEC-2000; 2000US-0251868.
PR 08-DEC-2000; 2000US-0251869.
PR 08-DEC-2000; 2000US-0251989.
PR 08-DEC-2000; 2000US-0251990.
PR 11-DEC-2000; 2000US-0254097.
PR 05-JAN-2001; 2001US-0259678.
XX
PA (HUMA-) HUMAN GENOME SCI INC.
XX
PI Rosen CA, Barash SC, Ruben SM;
XX
XX WPI, 2001-483426/52.
XX
XX Nucleic acids encoding human immune/hematopoietic antigen polypeptides,
PT useful for preventing, diagnosing and/or treating cancers and
PT metastasis -
XX

PR 13-OCT-2000; 2000US-0239937.
PR 20-OCT-2000; 2000US-0240960.
PR 20-OCT-2000; 2000US-0241321.
PR 20-OCT-2000; 2000US-0241785.
PR 20-OCT-2000; 2000US-0241786.
PR 20-OCT-2000; 2000US-0241787.
PR 20-OCT-2000; 2000US-0241808.
PR 20-OCT-2000; 2000US-0241809.
PR 20-OCT-2000; 2000US-0241826.
PR 01-NOV-2000; 2000US-0244617.
PR 08-NOV-2000; 2000US-0246474.
PR 08-NOV-2000; 2000US-0246475.
PR 08-NOV-2000; 2000US-0246476.
PR 08-NOV-2000; 2000US-0246477.
PR 08-NOV-2000; 2000US-0246478.
PR 08-NOV-2000; 2000US-0246523.
PR 08-NOV-2000; 2000US-0246524.
PR 08-NOV-2000; 2000US-0246525.
PR 08-NOV-2000; 2000US-0246526.
PR 08-NOV-2000; 2000US-0246527.
PR 08-NOV-2000; 2000US-0246528.
PR 08-NOV-2000; 2000US-0246532.
PR 08-NOV-2000; 2000US-0246609.
PR 08-NOV-2000; 2000US-0246610.
PR 08-NOV-2000; 2000US-0246611.
PR 17-NOV-2000; 2000US-0246613.
PR 17-NOV-2000; 2000US-0249207.
PR 17-NOV-2000; 2000US-0249208.
PR 17-NOV-2000; 2000US-0249209.
PR 17-NOV-2000; 2000US-0249210.
PR 17-NOV-2000; 2000US-0249211.
PR 17-NOV-2000; 2000US-0249212.
PR 17-NOV-2000; 2000US-0249213.
PR 17-NOV-2000; 2000US-0249214.
PR 17-NOV-2000; 2000US-0249215.
PR 17-NOV-2000; 2000US-0249216.
PR 17-NOV-2000; 2000US-0249217.
PR 17-NOV-2000; 2000US-0249218.
PR 17-NOV-2000; 2000US-0249219.
PR 17-NOV-2000; 2000US-0249244.
PR 17-NOV-2000; 2000US-0249245.
PR 17-NOV-2000; 2000US-0249264.
PR 17-NOV-2000; 2000US-0249265.
PR 17-NOV-2000; 2000US-0249297.
PR 17-NOV-2000; 2000US-0249299.
PR 17-NOV-2000; 2000US-0249300.
PR 01-DEC-2000; 2000US-0250160.
PR 01-DEC-2000; 2000US-0250391.
PR 05-DEC-2000; 2000US-0251030.
PR 05-DEC-2000; 2000US-0251988.
PR 05-DEC-2000; 2000US-0256719.
PR 06-DEC-2000; 2000US-0251479.
PR 08-DEC-2000; 2000US-0251856.
PR 08-DEC-2000; 2000US-0251868.
PR 08-DEC-2000; 2000US-0251869.
PR 08-DEC-2000; 2000US-0251989.
PR 08-DEC-2000; 2000US-0251990.
PR 11-DEC-2000; 2000US-0254097.
PR 05-JAN-2001; 2001US-0259678.
XX
XX
PA (HUMA-) HUMAN GENOME SCI INC.
XX
XX Rosen CA, Barash SC, Ruben SM;
XX WPI; 2001-457726/49.
XX
XX Isolated polypeptide for treating, preventing and prognosing disorders
PT related to the endocrine system including endocrine disorders,
PT reproductive disorders, and gastrointestinal disorders and also for
PT testing and detection e.g. diagnosis -
XX
XX Disclosure; SEQ ID NO 681; 558pp; English.
PS
XX The invention relates to cDNAs encoding novel human endocrine

CC antigens or a fragment having biological activity, a domain, an epitope,
CC full length protein, variant, allelic variant or a species homologue of
CC the cDNA/antigen. The DNAs and polypeptides are useful for preventing,
CC treating or ameliorating a medical condition when administered
CC (e.g. by gene therapy or antisense-therapy). Identifying mutations in
CC the genes coding for the antigens is useful for diagnosing a pathological
CC condition or a susceptibility to a pathological condition. The DNAs,
CC antigens and antibodies raised against the antigens useful for treating,
CC preventing and/or prognosing disorders related to the endocrine system
CC or hormone imbalance or reproductive disorders, cancers of endocrine
CC tissues, disorders of the pancreas (e.g. diabetes mellitus), the adrenal
CC glands (e.g. hirsutism), ovaries, the thyroid (e.g. hyperthyroidism), the
CC hypothalamus and testes (e.g. vanishing testes syndrome), many examples
CC of diseases and disorders are given in the specification. The present
CC sequence is genomic DNA fragment form a gene encoding an endocrine
CC antigen of the invention.
CC Note: The sequence data for this patent did not form part
CC of the printed specification, but was obtained in electronic
CC format directly from WIPO at
CC ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 17792 BP; 4372 A; 4626 C; 4413 G; 4381 T; 0 other;

Query Match 11.7%; Score 117.4; DB 22; Length 17792;
Best Local Similarity 81.1%; Pred. No. 4.6e-22;
Matches 150; Conservative 0; Mismatches 31; Indels 4; Gaps 1;

QY 1 TGGCAGCGCGCTGTAGTCCAGCTACTCAGGAGACTGAGCGAGAGATCCCTTGAACCC 60
DB 8122 TGGCAGGCACCTGTAGTCCAGCTACTCAGGAGGCTGAGGCGAGGAGATCCCTTGAACCT 8063

QY 61 GGGAGCGGAGGTTGCAGTGAGCAAGATCGCTGCTACTGCTCCAGCTCGGCGACA--- 117
DB 8062 GGGAGCGGAGGTTGCAGTGAGCTGAGATGTCACCTGCTCCAGCTCGGCGACAAG 8003

QY 118 -GAGTTCGCTTCCAAAGAAAAAATAATTAATAAAGATAAATCCGCGCTGC 176
DB 8002 CGAGACTCCGCTCTCAAAAAAATAATAATAATAATAAATAAATTCGCGGCG 7943

QY 177 GCGGT 181
DB 7942 ATGCT 7938

RESULT 8
AAS36099/C
ID AAS36099 standard; DNA; 17792 BP.
XX
AC AAS36099;
DT 17-DEC-2001 (first entry)
DE Human cardiovascular system antigen genomic DNA SEQ ID No 1599.
XX
KW Cardiovascular system antigen; human; mouse; rabbit; goat; horse; cat;
KW chicken; sheep; immunosuppressive; antiarthritic; vasotropic; dog;
KW antirheumatic; antiproliferative; cytostatic; cardiac; neuroprotective;
KW cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer;
KW ophthalmological; vulnery; gene therapy; autoimmune disease; neoplasm;
KW hyperproliferative disorder; breast; liver; cardiovascular disorder;
KW cerebrovascular disorder; nervous system disorder; bacterial infection;
KW fungal infection; viral infection; ocular disorder; endocrine disorder;
KW gastrointestinal disorder; renal disorder; respiratory disorder;
KW wound healing; skin aging; organ transplantation; tissue regeneration;
KW anti-infertility.
XX
OS Homo sapiens.
XX
PN WO200155321-A2.
XX
PD 02-AUG-2001.
XX
PF 17-JAN-2001; 2001WO-US01340.

PT cardiovascular system -
PS Claim 1; SEQ ID No 1599; 674pp; English.
XX
XX Sequences AAS35741-AAS36942 represent genomic DNA molecules, which encode
CC the cardiovascular system antigen polypeptides of the invention.
CC Cardiovascular system antigens and their associated polynucleotides are
CC useful in the diagnosis, treatment and prevention of various types of
CC disorders in e.g. humans, mice, rabbits, goats, horses, cats, dogs,
CC chickens or sheep. A pathological condition can be determined by
CC detecting the presence or absence of a mutation in a cardiovascular
CC system antigen polynucleotide. The treatable disorders include autoimmune
CC diseases such as rheumatoid arthritis, hyperproliferative disorders such
CC as neoplasms of the breast or liver, cardiovascular disorders such as
CC cardiac arrest, cerebrovascular disorders such as cerebral ischaemia,
CC nervous system disorders such as Alzheimer's disease, infections caused
CC by bacteria, viruses and fungi, ocular disorders such as corneal
CC infection, endocrine disorders such as premature labour and infertility,
CC gastrointestinal disorders such as Crohn's disease, renal disorders such
CC as glomerulonephritis and respiratory disorders such as asthma and
CC pleurisy. The polypeptides can also be used to aid wound healing, to
CC prevent skin aging due to sunburn, to maintain organs before
CC transplantation, to regenerate tissues and in chemotaxis.
CC Note: The sequence data for this patent did not form part of the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences.

Query Match 11.7%; Score 117.4; DB 22; Length 17792;
Best Local Similarity 81.1%; Pred. No. 4.6e-22;
Matches 150; Conservative 0; Mismatches 31; Indels 4; Gaps 1;
XX
XX 1 TGGCAGCGCGCTGTAGTCCCGAGCTACTCAGGAGCTAGGCGAGGAGATCGCTTGAACCC 60
Db 8122 TGGCAGCGCGCTGTAGTCCCGAGCTACTCAGGAGCTAGGCGAGGAGATCGCTTGAACCT 8063
XX
XX 61 GGGAGCGGAGGTTGCAGTGGAGCAAGATCGCGTCACTGCAGCTCCAGCTGGCGGACA --- 117
Db 8062 GGGAGCGGAGGTTGCAGTGGAGCAAGATCGCGTCACTGCAGCTCCAGCTGGCGGACAAG 8003
XX
XX 118 -GACGTCCTCGTTTCAAAAGAAAAAATAATTAATAAAAGATAAATCGCGGCTGC 176
Db 8002 CGAGACTCCGCTCAAAAAAATAATAATAATAATAATAATAATAATAATAATAATAATAATA 7943
XX
XX 177 GCGGT 181
Db 7942 ATGGT 7938
XX
XX
XX RESULT 9
XX AAX83003
XX ID AAX83003 standard; DNA; 87350 BP.
XX AC AAX83003;
XX
XX 31-AUG-1999 (first entry)
XX Human WRN genomic sequence.
XX
XX Human; WRN; Werner's syndrome; detection; diagnosis; autosomal;
XX recessive disorder; phenotype: ss.
XX Homo sapiens.
XX
XX WO9724435-A1.
XX
XX 10-JUL-1997.
XX
XX 30-DEC-1996; 96WO-US20785.
XX
XX 12-APR-1996; 96US-0632175.
XX 29-DEC-1995; 95US-0009409.
XX 29-DEC-1995; 95US-0580539.
XX

PR 30-JAN-1996; 96US-0010835.
PR 30-JAN-1996; 96US-0594242.
XX
XX (DARW-) DARWIN MOLECULAR CORP.
XX (OSHI/) OSHIMA J.
XX Fu Y, Mulligan J, Oshima J, Schellenberg GD, Yu C;
XX WPI; 1997-363671/33.
XX Isolated nucleic acid molecule encoding the WRN gene product -
XX useful for detection and treatment of Werner's syndrome, and related
XX diseases
XX
XX Claim 1; Fig 5A-U; 153pp; English.
XX
XX This sequence represents the genomic region containing the coding
XX sequence for the human WRN gene which encodes a protein related to
XX Werner's syndrome. The products can be used for the detection and
XX treatment of Werner's syndrome (WS), an autosomal recessive disorder
XX with a complex phenotype, as well as related diseases.
XX
XX Sequence 87350 BP; 25621 A; 16221 C; 17012 G; 28450 T; 46 other;
XX
XX Query Match 11.7%; Score 117; DB 18; Length 87350;
XX Best Local Similarity 81.8%; Pred. No. 1.3e-21;
XX Matches 135; Conservative 0; Mismatches 30; Indels 0; Gaps 0;
XX
XX 1 TGGCAGCGCGCTGTAGTCCCGAGCTACTCAGGAGCTAGGCGAGGAGATCGCTTGAACCC 60
Db 10250 TGGCAGCGCGCTGTAGTCCCGAGCTACTTGGGAGGCTAGGCGAGGAGATCGCTTGAACCT 10309
XX
XX 61 GGGAGCGGAGGTTGCAGTGGAGCAAGATCGCGTCACTGCAGCTCCAGCTGGCGGACAGAC 120
Db 10310 GGGAGCGGAGGTTGCAGTGGAGCAAGATCGCGTCACTGCAGCTCCAGCTGGCGGACAGAC 10369
XX
XX 121 GTTCCGTTTCAAAAGAAAAAATAATTAATAAAAGATAAATAA 165
Db 10370 CGACACTCTTCTCAAAAAAATAAATAAATAAATAAATAA 10414
XX
XX
XX RESULT 10
XX AAK62290/c
XX ID AAK62290 standard; cDNA; 1039 BP.
XX AC AAK62290;
XX
XX 06-NOV-2001 (first entry)
XX Human immune/haematopoietic antigen encoding cDNA SEQ ID NO:7350.
XX
XX Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
XX cytostatic; gene therapy; vaccine; metastasis; ss.
XX Homo sapiens.
XX
XX WO200157182-A2.
XX
XX 09-AUG-2001.
XX
XX 17-JAN-2001; 2001WO-US01354.
XX
XX 31-JAN-2000; 2000US-0179065.
XX 04-FEB-2000; 2000US-0180628.
XX 24-FEB-2000; 2000US-0184664.
XX 02-MAR-2000; 2000US-0186350.
XX 16-MAR-2000; 2000US-0189874.
XX 17-MAR-2000; 2000US-0190076.
XX 18-APR-2000; 2000US-0198123.
XX 19-MAY-2000; 2000US-0205515.
XX 07-JUN-2000; 2000US-0209467.
XX 28-JUN-2000; 2000US-0214886.
XX 30-JUN-2000; 2000US-0215135.
XX

PR 07-JUL-2000; 2000US-0216647.
PR 07-JUL-2000; 2000US-0216880.
PR 11-JUL-2000; 2000US-0217487.
PR 11-JUL-2000; 2000US-0217496.
PR 14-JUL-2000; 2000US-0218290.
PR 26-JUL-2000; 2000US-0220963.
PR 26-JUL-2000; 2000US-0220964.
PR 14-AUG-2000; 2000US-0224518.
PR 14-AUG-2000; 2000US-0224519.
PR 14-AUG-2000; 2000US-0225213.
PR 14-AUG-2000; 2000US-0225214.
PR 14-AUG-2000; 2000US-0225266.
PR 14-AUG-2000; 2000US-0225267.
PR 14-AUG-2000; 2000US-0225268.
PR 14-AUG-2000; 2000US-0225270.
PR 14-AUG-2000; 2000US-0225447.
PR 14-AUG-2000; 2000US-0225757.
PR 14-AUG-2000; 2000US-0225758.
PR 14-AUG-2000; 2000US-0225759.
PR 18-AUG-2000; 2000US-0226279.
PR 22-AUG-2000; 2000US-0226681.
PR 22-AUG-2000; 2000US-0226688.
PR 22-AUG-2000; 2000US-0227182.
PR 23-AUG-2000; 2000US-0227009.
PR 30-AUG-2000; 2000US-0228924.
PR 01-SEP-2000; 2000US-0229287.
PR 01-SEP-2000; 2000US-0229343.
PR 01-SEP-2000; 2000US-0229344.
PR 01-SEP-2000; 2000US-0229345.
PR 05-SEP-2000; 2000US-0229509.
PR 05-SEP-2000; 2000US-0229513.
PR 06-SEP-2000; 2000US-0230437.
PR 06-SEP-2000; 2000US-0230438.
PR 08-SEP-2000; 2000US-0231242.
PR 08-SEP-2000; 2000US-0231243.
PR 08-SEP-2000; 2000US-0231244.
PR 08-SEP-2000; 2000US-0231413.
PR 08-SEP-2000; 2000US-0231414.
PR 08-SEP-2000; 2000US-0232080.
PR 08-SEP-2000; 2000US-0232081.
PR 12-SEP-2000; 2000US-0231968.
PR 14-SEP-2000; 2000US-0232397.
PR 14-SEP-2000; 2000US-0232398.
PR 14-SEP-2000; 2000US-0232399.
PR 14-SEP-2000; 2000US-0232400.
PR 14-SEP-2000; 2000US-0232401.
PR 14-SEP-2000; 2000US-0233063.
PR 14-SEP-2000; 2000US-0233064.
PR 14-SEP-2000; 2000US-0233065.
PR 21-SEP-2000; 2000US-0234223.
PR 21-SEP-2000; 2000US-0234274.
PR 25-SEP-2000; 2000US-0234997.
PR 25-SEP-2000; 2000US-0234998.
PR 26-SEP-2000; 2000US-0235484.
PR 27-SEP-2000; 2000US-0235834.
PR 27-SEP-2000; 2000US-0235836.
PR 29-SEP-2000; 2000US-0236327.
PR 29-SEP-2000; 2000US-0236367.
PR 29-SEP-2000; 2000US-0236368.
PR 29-SEP-2000; 2000US-0236369.
PR 29-SEP-2000; 2000US-0236370.
PR 02-OCT-2000; 2000US-0236802.
PR 02-OCT-2000; 2000US-0237037.
PR 02-OCT-2000; 2000US-0237038.
PR 02-OCT-2000; 2000US-0237039.
PR 02-OCT-2000; 2000US-0237040.
PR 13-OCT-2000; 2000US-0239935.
PR 13-OCT-2000; 2000US-0239937.
PR 20-OCT-2000; 2000US-0240960.
PR 20-OCT-2000; 2000US-0241221.
PR 20-OCT-2000; 2000US-0241785.
PR 20-OCT-2000; 2000US-0241786.
PR 20-OCT-2000; 2000US-0241787.
PR 20-OCT-2000; 2000US-0241808.
PR 20-OCT-2000; 2000US-0241809.
PR 20-OCT-2000; 2000US-0241826.
PR 01-NOV-2000; 2000US-0244617.
PR 08-NOV-2000; 2000US-0246474.
PR 08-NOV-2000; 2000US-0246475.
PR 08-NOV-2000; 2000US-0246476.
PR 08-NOV-2000; 2000US-0246477.
PR 08-NOV-2000; 2000US-0246478.
PR 08-NOV-2000; 2000US-0246523.
PR 08-NOV-2000; 2000US-0246524.
PR 08-NOV-2000; 2000US-0246525.
PR 08-NOV-2000; 2000US-0246526.
PR 08-NOV-2000; 2000US-0246527.
PR 08-NOV-2000; 2000US-0246528.
PR 08-NOV-2000; 2000US-0246532.
PR 08-NOV-2000; 2000US-0246609.
PR 08-NOV-2000; 2000US-0246610.
PR 08-NOV-2000; 2000US-0246611.
PR 08-NOV-2000; 2000US-0246613.
PR 17-NOV-2000; 2000US-0249207.
PR 17-NOV-2000; 2000US-0249208.
PR 17-NOV-2000; 2000US-0249209.
PR 17-NOV-2000; 2000US-0249210.
PR 17-NOV-2000; 2000US-0249211.
PR 17-NOV-2000; 2000US-0249212.
PR 17-NOV-2000; 2000US-0249213.
PR 17-NOV-2000; 2000US-0249214.
PR 17-NOV-2000; 2000US-0249215.
PR 17-NOV-2000; 2000US-0249216.
PR 17-NOV-2000; 2000US-0249217.
PR 17-NOV-2000; 2000US-0249218.
PR 17-NOV-2000; 2000US-0249244.
PR 17-NOV-2000; 2000US-0249245.
PR 17-NOV-2000; 2000US-0249264.
PR 17-NOV-2000; 2000US-0249265.
PR 17-NOV-2000; 2000US-0249297.
PR 17-NOV-2000; 2000US-0249299.
PR 17-NOV-2000; 2000US-0249300.
PR 01-DEC-2000; 2000US-0250160.
PR 01-DEC-2000; 2000US-0250391.
PR 05-DEC-2000; 2000US-0251030.
PR 05-DEC-2000; 2000US-0251988.
PR 06-DEC-2000; 2000US-0256719.
PR 08-DEC-2000; 2000US-0251479.
PR 08-DEC-2000; 2000US-0251856.
PR 08-DEC-2000; 2000US-0251868.
PR 08-DEC-2000; 2000US-0251869.
PR 08-DEC-2000; 2000US-0251989.
PR 08-DEC-2000; 2000US-0251990.
PR 11-DEC-2000; 2000US-0254097.
PR 05-JAN-2001; 2001US-0259678.
XX
XX
PA (HUMA-) HUMAN GENOME SCI INC.
XX
XX Rosen CA, Barash SC, Ruben SM;
XX
XX WPI: 2001-483426/52.
DR P-FSDB; AAM89509.
XX
XX Nucleic acids encoding human immune/hematopoietic antigen polypeptides,
PT useful for preventing, diagnosing and/or treating cancers and
PT metastasis -
XX
XX Claim 1; SEQ ID NO 7350; 3071pp + Sequence Listing; English.
PS
XX
XX AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I)
CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic
CC activity, and can be used in gene therapy and vaccine production. (I)
CC proteins and polynucleotides may be used in the prevention, diagnosis and
CC treatment of diseases associated with inappropriate (I) expression. For
CC example, they may be used to treat disorders associated with decreased
CC expression by rectifying mutations or deletions in a patient's genome

PR 18-AUG-2000; 2000US-0226279.
PR 22-AUG-2000; 2000US-0226681.
PR 22-AUG-2000; 2000US-0226868.
PR 23-AUG-2000; 2000US-0227182.
PR 23-AUG-2000; 2000US-0227009.
PR 30-AUG-2000; 2000US-0228924.
PR 01-SEP-2000; 2000US-0229287.
PR 01-SEP-2000; 2000US-0229343.
PR 01-SEP-2000; 2000US-0229344.
PR 01-SEP-2000; 2000US-0229345.
PR 05-SEP-2000; 2000US-0229509.
PR 05-SEP-2000; 2000US-0229513.
PR 06-SEP-2000; 2000US-0230437.
PR 06-SEP-2000; 2000US-0230438.
PR 08-SEP-2000; 2000US-0231242.
PR 08-SEP-2000; 2000US-0231243.
PR 08-SEP-2000; 2000US-0231244.
PR 08-SEP-2000; 2000US-0231413.
PR 08-SEP-2000; 2000US-0231414.
PR 08-SEP-2000; 2000US-0231418.
PR 08-SEP-2000; 2000US-0232081.
PR 12-SEP-2000; 2000US-0231968.
PR 14-SEP-2000; 2000US-0232397.
PR 14-SEP-2000; 2000US-0232398.
PR 14-SEP-2000; 2000US-0232399.
PR 14-SEP-2000; 2000US-0232400.
PR 14-SEP-2000; 2000US-0232401.
PR 14-SEP-2000; 2000US-0233063.
PR 14-SEP-2000; 2000US-0233064.
PR 14-SEP-2000; 2000US-0233065.
PR 21-SEP-2000; 2000US-0234223.
PR 21-SEP-2000; 2000US-0234274.
PR 25-SEP-2000; 2000US-0234997.
PR 25-SEP-2000; 2000US-0234998.
PR 26-SEP-2000; 2000US-0235484.
PR 27-SEP-2000; 2000US-0235834.
PR 27-SEP-2000; 2000US-0235836.
PR 29-SEP-2000; 2000US-0236327.
PR 29-SEP-2000; 2000US-0236367.
PR 29-SEP-2000; 2000US-0236368.
PR 29-SEP-2000; 2000US-0236369.
PR 29-SEP-2000; 2000US-0236370.
PR 02-OCT-2000; 2000US-0236802.
PR 02-OCT-2000; 2000US-0237037.
PR 02-OCT-2000; 2000US-0237038.
PR 02-OCT-2000; 2000US-0237039.
PR 02-OCT-2000; 2000US-0237040.
PR 13-OCT-2000; 2000US-0239935.
PR 13-OCT-2000; 2000US-0239937.
PR 20-OCT-2000; 2000US-0240960.
PR 20-OCT-2000; 2000US-0241221.
PR 20-OCT-2000; 2000US-0241785.
PR 20-OCT-2000; 2000US-0241786.
PR 20-OCT-2000; 2000US-0241787.
PR 20-OCT-2000; 2000US-0241808.
PR 20-OCT-2000; 2000US-0241809.
PR 20-OCT-2000; 2000US-0241826.
PR 01-NOV-2000; 2000US-0244617.
PR 01-NOV-2000; 2000US-0244674.
PR 08-NOV-2000; 2000US-0246475.
PR 08-NOV-2000; 2000US-0246476.
PR 08-NOV-2000; 2000US-0246477.
PR 08-NOV-2000; 2000US-0246478.
PR 08-NOV-2000; 2000US-0246523.
PR 08-NOV-2000; 2000US-0246524.
PR 08-NOV-2000; 2000US-0246525.
PR 08-NOV-2000; 2000US-0246526.
PR 08-NOV-2000; 2000US-0246527.
PR 08-NOV-2000; 2000US-0246528.
PR 08-NOV-2000; 2000US-0246532.
PR 08-NOV-2000; 2000US-0246609.
PR 08-NOV-2000; 2000US-0246610.
PR 08-NOV-2000; 2000US-0246611.

PR 08-NOV-2000; 2000US-0246613.
PR 17-NOV-2000; 2000US-0249207.
PR 17-NOV-2000; 2000US-0249208.
PR 17-NOV-2000; 2000US-0249209.
PR 17-NOV-2000; 2000US-0249210.
PR 17-NOV-2000; 2000US-0249211.
PR 17-NOV-2000; 2000US-0249212.
PR 17-NOV-2000; 2000US-0249213.
PR 17-NOV-2000; 2000US-0249214.
PR 17-NOV-2000; 2000US-0249215.
PR 17-NOV-2000; 2000US-0249216.
PR 17-NOV-2000; 2000US-0249217.
PR 17-NOV-2000; 2000US-0249218.
PR 17-NOV-2000; 2000US-0249244.
PR 17-NOV-2000; 2000US-0249245.
PR 17-NOV-2000; 2000US-0249264.
PR 17-NOV-2000; 2000US-0249265.
PR 17-NOV-2000; 2000US-0249297.
PR 17-NOV-2000; 2000US-0249299.
PR 17-NOV-2000; 2000US-0249300.
PR 01-DEC-2000; 2000US-0250160.
PR 01-DEC-2000; 2000US-0250391.
PR 05-DEC-2000; 2000US-0251030.
PR 05-DEC-2000; 2000US-0251988.
PR 05-DEC-2000; 2000US-0256719.
PR 06-DEC-2000; 2000US-0251479.
PR 08-DEC-2000; 2000US-0251856.
PR 08-DEC-2000; 2000US-0251868.
PR 08-DEC-2000; 2000US-0251869.
PR 08-DEC-2000; 2000US-0251989.
PR 08-DEC-2000; 2000US-0251990.
PR 11-DEC-2000; 2000US-0254097.
PR 05-JAN-2001; 2001US-0259678.
XX
XX
PA (HUMA-) HUMAN GENOME SCI INC.
XX
XX Rosen CA, Barash SC, Ruben SM;
PI
XX
XX WPI; 2001-483426/52.
DR P-PSDB; AAM91778.
XX
PT Nucleic acids encoding human immune/hematopoietic antigen polypeptides,
PT useful for preventing, diagnosing and/or treating cancers and
PT metastasis -
XX
PS Claim 1; SEQ ID NO 9619; 3071pp + Sequence Listing; English.
XX
CC AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I)
CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic
CC activity, and can be used in gene therapy and vaccine production. (I)
CC proteins and polynucleotides may be used in the prevention, diagnosis and
CC treatment of diseases associated with inappropriate (I) expression. For
CC example, they may be used to treat disorders associated with decreased
CC expression by rectifying mutations or deletions in a patient's genome
CC that affect the activity of (I) by expressing inactive proteins or to
CC supplement the patients own production of (I). Additionally, (I)
CC polynucleotides may be used to produce the secreted (I), by inserting
CC the nucleic acids into a host cell and culturing the cell to express the
CC protein. (I) proteins and polynucleotides may be used to prevent,
CC diagnose and treat immune/haematopoietic-related diseases, especially
CC cancers and cancer metastases of haematopoietic-derived cells. AAK64703
CC to AAK87694 represent human immune/haematopoietic antigen genomic
CC sequences from the present invention. AAK54942 to AAK54950 and AAM82169
CC represent sequences used in the exemplification of the present invention.
XX
SQ Sequence 666 BP; 139 A; 177 C; 146 G; 199 T; 5 other;

Query Match 11.5%; Score 115.4; DB 22; Length 666;
Best Local Similarity 84.0%; Pred. No. 3.3e-22;
Matches 142; Conservative 0; Mismatches 23; Indels 4; Gaps 1;
QY 1 TGGCACGCGCTGTAGTCCCGAGCTACTCAGGAGACTAGGAGGAGGAGTCCGTTGAACCC 60
||||| ||||||||| ||||||| |||| ||||||||| ||||||| ||||||| ||||||| |||||||

PR	26-JUL-2000;	2000US-0220954.	PR	08-NOV-2000;	2000US-0246476.
PR	14-AUG-2000;	2000US-0220954.	PR	08-NOV-2000;	2000US-0246477.
PR	14-AUG-2000;	2000US-0224518.	PR	08-NOV-2000;	2000US-0246478.
PR	14-AUG-2000;	2000US-0224519.	PR	08-NOV-2000;	2000US-0246523.
PR	14-AUG-2000;	2000US-0225213.	PR	08-NOV-2000;	2000US-0246524.
PR	14-AUG-2000;	2000US-0225214.	PR	08-NOV-2000;	2000US-0246525.
PR	14-AUG-2000;	2000US-0225266.	PR	08-NOV-2000;	2000US-0246526.
PR	14-AUG-2000;	2000US-0225287.	PR	08-NOV-2000;	2000US-0246527.
PR	14-AUG-2000;	2000US-0225288.	PR	08-NOV-2000;	2000US-0246528.
PR	14-AUG-2000;	2000US-0225270.	PR	08-NOV-2000;	2000US-0246532.
PR	14-AUG-2000;	2000US-0225247.	PR	08-NOV-2000;	2000US-0246609.
PR	14-AUG-2000;	2000US-0225757.	PR	08-NOV-2000;	2000US-0246610.
PR	14-AUG-2000;	2000US-0225758.	PR	08-NOV-2000;	2000US-0246611.
PR	14-AUG-2000;	2000US-0225759.	PR	08-NOV-2000;	2000US-0246613.
PR	18-AUG-2000;	2000US-0226279.	PR	17-NOV-2000;	2000US-0249207.
PR	22-AUG-2000;	2000US-0226681.	PR	17-NOV-2000;	2000US-0249208.
PR	22-AUG-2000;	2000US-0226868.	PR	17-NOV-2000;	2000US-0249209.
PR	22-AUG-2000;	2000US-0227182.	PR	17-NOV-2000;	2000US-0249210.
PR	23-AUG-2000;	2000US-0227009.	PR	17-NOV-2000;	2000US-0249211.
PR	30-AUG-2000;	2000US-0228924.	PR	17-NOV-2000;	2000US-0249212.
PR	03-SEP-2000;	2000US-0229287.	PR	17-NOV-2000;	2000US-0249213.
PR	01-SEP-2000;	2000US-0229343.	PR	17-NOV-2000;	2000US-0249214.
PR	01-SEP-2000;	2000US-0229344.	PR	17-NOV-2000;	2000US-0249215.
PR	01-SEP-2000;	2000US-0229345.	PR	17-NOV-2000;	2000US-0249216.
PR	05-SEP-2000;	2000US-0229509.	PR	17-NOV-2000;	2000US-0249217.
PR	05-SEP-2000;	2000US-0229513.	PR	17-NOV-2000;	2000US-0249218.
PR	06-SEP-2000;	2000US-0230437.	PR	17-NOV-2000;	2000US-0249219.
PR	06-SEP-2000;	2000US-0230438.	PR	17-NOV-2000;	2000US-0249244.
PR	08-SEP-2000;	2000US-0231242.	PR	17-NOV-2000;	2000US-0249245.
PR	08-SEP-2000;	2000US-0231243.	PR	17-NOV-2000;	2000US-0249246.
PR	08-SEP-2000;	2000US-0231244.	PR	17-NOV-2000;	2000US-0249265.
PR	08-SEP-2000;	2000US-0231413.	PR	17-NOV-2000;	2000US-0249297.
PR	08-SEP-2000;	2000US-0231414.	PR	17-NOV-2000;	2000US-0249299.
PR	08-SEP-2000;	2000US-0232080.	PR	01-DEC-2000;	2000US-0250391.
PR	08-SEP-2000;	2000US-0232081.	PR	01-DEC-2000;	2000US-0251160.
PR	12-SEP-2000;	2000US-0231968.	PR	05-DEC-2000;	2000US-0251030.
PR	14-SEP-2000;	2000US-0232397.	PR	05-DEC-2000;	2000US-0251988.
PR	14-SEP-2000;	2000US-0232398.	PR	05-DEC-2000;	2000US-0256719.
PR	14-SEP-2000;	2000US-0232399.	PR	06-DEC-2000;	2000US-0251479.
PR	14-SEP-2000;	2000US-0232400.	PR	08-DEC-2000;	2000US-0251856.
PR	14-SEP-2000;	2000US-0232401.	PR	08-DEC-2000;	2000US-0251868.
PR	14-SEP-2000;	2000US-0233063.	PR	08-DEC-2000;	2000US-0251869.
PR	14-SEP-2000;	2000US-0233084.	PR	08-DEC-2000;	2000US-0251989.
PR	14-SEP-2000;	2000US-0233085.	PR	08-DEC-2000;	2000US-0251990.
PR	21-SEP-2000;	2000US-0234223.	PR	11-DEC-2000;	2000US-0254097.
PR	21-SEP-2000;	2000US-0234274.	PR	05-JAN-2001;	2001US-0259678.
PR	25-SEP-2000;	2000US-0234997.	PR		
PR	25-SEP-2000;	2000US-0234998.	XX		
PR	26-SEP-2000;	2000US-0235484.	PA		
PR	27-SEP-2000;	2000US-0235834.			
PR	27-SEP-2000;	2000US-0235836.	PI		
PR	29-SEP-2000;	2000US-0236327.	PI	</	

PR 08-NOV-2000; 2000US-0246610.
PR 08-NOV-2000; 2000US-0246611.
PR 08-NOV-2000; 2000US-0246613.
PR 17-NOV-2000; 2000US-0249207.
PR 17-NOV-2000; 2000US-0249208.
PR 17-NOV-2000; 2000US-0249209.
PR 17-NOV-2000; 2000US-0249210.
PR 17-NOV-2000; 2000US-0249211.
PR 17-NOV-2000; 2000US-0249212.
PR 17-NOV-2000; 2000US-0249213.
PR 17-NOV-2000; 2000US-0249214.
PR 17-NOV-2000; 2000US-0249215.
PR 17-NOV-2000; 2000US-0249216.
PR 17-NOV-2000; 2000US-0249217.
PR 17-NOV-2000; 2000US-0249218.
PR 17-NOV-2000; 2000US-0249244.
PR 17-NOV-2000; 2000US-0249245.
PR 17-NOV-2000; 2000US-0249284.
PR 17-NOV-2000; 2000US-0249285.
PR 17-NOV-2000; 2000US-0249297.
PR 17-NOV-2000; 2000US-0249299.
PR 17-NOV-2000; 2000US-0249300.
PR 01-DEC-2000; 2000US-0250160.
PR 01-DEC-2000; 2000US-0250391.
PR 05-DEC-2000; 2000US-0251030.
PR 05-DEC-2000; 2000US-0251988.
PR 05-DEC-2000; 2000US-0256719.
PR 06-DEC-2000; 2000US-0251479.
PR 08-DEC-2000; 2000US-0251856.
PR 08-DEC-2000; 2000US-0251868.
PR 08-DEC-2000; 2000US-0251869.
PR 08-DEC-2000; 2000US-0251989.
PR 08-DEC-2000; 2000US-0251990.
PR 11-DEC-2000; 2000US-0254097.
PR 05-JAN-2001; 2001US-0259678.

(HUMA-) HUMAN GENOME SCI INC.

Rosen CA, Barash SC, Ruben SM;

WPI; 2001-483426/52.

Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis -

Disclosure; SEQ ID NO 21743; 3071pp + Sequence Listing; English.

AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I) by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK54942 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention.

Sequence 30620 BP; 7407 A; 7216 C; 7192 G; 8805 T; 0 other;

Query Match 11.4%; Score 114; DB 22; Length 30620;

Best Local Similarity 74.2%; Pred. No. 5.4e-21;

Matches 144; Conservative 0; Mismatches 50; Indels 0; Gaps 0;

QY 1 TGGCAGCGCGCTGTAGTCCAGCTACTCAGGAGACTGAGGAGGAGAGAAATCGTTGAACCC 60

Db 18324 TGGCAGCGCGCTGTAAATCCAGCTACTCGGAGGCTGAGGAGAGAAATCACTTGAACCC 18265
QY 61 GGGAGACGGAGGTTGCACTGAGCCAAAGATCGCGTCACCTCCAGCTCGCGACAGAC 120
Db 18264 GGGAGGACAGGTTGCGGTGAGCCCTAGATCATGTCTCGCACTCCAGCTCGGGGACAAG 18205
QY 121 GTTCCGTTTCAAAAGAAAAAATAATATTAATAAAAAAGAAATAAAATCGCGGCTCGCGG 180
Db 18204 TGTGAGACTTCATCTCAAAAAAAGAAAAAAGAAAAAAGCGCGCGGTGCGG 18145
QY 181 TGACATCAGTCTCT 194
Db 18144 TGGCTCAGCTTCT 18131

Search completed: December 24, 2002, 17:19:36
Job time : 270.952 secs

THIS PAGE BLANK (USPTO)

Db 5506 CAGCCCTGACTTGG-----GGAGGGGTGAAAACAGAGACCCACCAGGGTAGGACATTTCAA 5452
QY 475 CTGGCTCCATCCTCTGCATCTTTAGATTATTTGGGACAGTTTGATACAGAGAAGGAGGA 534
II III II II III III III III III III III III III III III III III III III
Db 5451 CTCCTACCTCTCCTTAGATTTCATGGGACATGGGACATTTGGACACAGAGAAG--GGG 5394
QY 535 GACCCATCCCAATGAGGGTTTGATTAGATGAATATATCAATGATAATTCCTAGAGGA 594
III III III III III III III III III III III III III III III III III III
Db 5393 CTCCTCCCATGAGGGGTGAGGTAGAGGTTTAAATCCAGGACCACCTCTCTG----- 5339
QY 595 GGGACCTTTTATAATCACTCTGAGACAGGTTGGAGCTACATGGGATTTGGAGGGGAGGG 654
Db 5338 GGGAGTGGCTGAGCCCACTTTGAGAACAGGAAGGC-CTGGATGGGAGTAGAGGGGGGGA 5280
QY 655 TGGAGCCCTTTAAAAGAAAGCCCGAGAGACTGCCCCCTGCTCTCTCTCCCAACAAGT 714
II III II III III III III III III III III III III III III III III III
Db 5279 AGGGACCCC--AAGAACAAAGACCAGAGACTGGCCCTGTCTGCCCGGGGGCTTGT 5222
QY 715 TCCATTATTAATCTTCCACCCAGGAGCTGTGAGAAATCCTGCCCTTCTCTCCAGATCAA 774
Db 5221 CACCGCCCTGTCTTCC-----CTCTGGCCCGCAGGAGTTTGA 5185
QY 775 AGTCCTTCAGGAATGCAACTACTTCACT-GACAAGAGATATATATCTTCTGACAGA 833
Db 5184 GGTGATGCCCGAGATCAAGCTGCTGCACTGGCGCTGCAACAACTACAGCATGCGGCCAGA 5125
QY 834 GGAGGAATTTGGGGTTTGGTCCAGTCCATGAAGTGGCACAGTACAGATAAAAGCTGAGA 893
Db 5124 TGAGCAATTTGGGGCTGTTCGGGGCGGTGGAGCGGCTCAGCGAGACTGAGAGGTGAGG 5065
QY 894 GCTTAGGAG-AATTAGCGAGGGTAGAAGAACACTCTGTCTTTGTGACCACTTCAGAGAGC 952
Db 5064 CCGGGGAGCAAAATGGGACAGTGGGGCAGGCTGTCTTAGGGGCCAGCTCCAGAGGGC 5005
QY 953 CTGGGGCCATGCTTCCTGCTCAACATAGGCGCTGCTGCATGGTGA 999
Db 5004 CAGCAGCTATGAC-CCATGGTCAAGCTCAGCCCTGTTGCTGTGAGGA 4959

RESULT 3
ABL68560
ID ABL68560 standard; DNA; 267156 BP.
XX AC ABL68560;
XX DT 15-MAY-2002 (first entry)
XX DE Kidney cancer related gene sequence SEQ ID NO:6897.
XX KW Human; cancer; colon; breast; ovary; oesophagus; kidney; thyroid;
KW stomach; lung; prostate; pancreas; carcinoma; antitumour; cancerous;
KW cytostatic; gene therapy; antineoplastic; Wilms' tumour; adenocarcinoma;
KW gene; ds.
XX OS Homo sapiens.
XX PN WO200194629-A2.
XX PD 13-DEC-2001.
XX PF 30-MAY-2001; 2001WO-US10838.
XX PR 05-JUN-2000; 2000US-209473P.
PR 05-JUN-2000; 2000US-209531P.
PR 18-SEP-2000; 2000US-233133P.
PR 18-SEP-2000; 2000US-233617P.
PR 20-SEP-2000; 2000US-234009P.
PR 20-SEP-2000; 2000US-234034P.
PR 20-SEP-2000; 2000US-234052P.
PR 22-SEP-2000; 2000US-234509P.
PR 22-SEP-2000; 2000US-234567P.
PR 25-SEP-2000; 2000US-234923P.
PR 25-SEP-2000; 2000US-234924P.

PR 25-SEP-2000; 2000US-235077P.
PR 25-SEP-2000; 2000US-235082P.
PR 25-SEP-2000; 2000US-235134P.
PR 26-SEP-2000; 2000US-235280P.
PR 26-SEP-2000; 2000US-235637P.
PR 26-SEP-2000; 2000US-235638P.
PR 27-SEP-2000; 2000US-235711P.
PR 27-SEP-2000; 2000US-235720P.
PR 27-SEP-2000; 2000US-235840P.
PR 27-SEP-2000; 2000US-235863P.
PR 28-SEP-2000; 2000US-236028P.
PR 28-SEP-2000; 2000US-236032P.
PR 28-SEP-2000; 2000US-236033P.
PR 28-SEP-2000; 2000US-236034P.
PR 28-SEP-2000; 2000US-236109P.
PR 29-SEP-2000; 2000US-236111P.
PR 29-SEP-2000; 2000US-236842P.
PR 29-SEP-2000; 2000US-236891P.
PR 02-OCT-2000; 2000US-237172P.
PR 02-OCT-2000; 2000US-237173P.
PR 02-OCT-2000; 2000US-237278P.
PR 02-OCT-2000; 2000US-237294P.
PR 02-OCT-2000; 2000US-237295P.
PR 02-OCT-2000; 2000US-237316P.
PR 03-OCT-2000; 2000US-237425P.
PR 03-OCT-2000; 2000US-237598P.
PR 03-OCT-2000; 2000US-237604P.
PR 03-OCT-2000; 2000US-237606P.
PR 03-OCT-2000; 2000US-237608P.
PR 01-NOV-2000; 2000US-244867P.
PR 01-NOV-2000; 2000US-245084P.
PA (AVAL-) AVALON PHARM.
XX Young PE, Augustus M, Carter KC, Ebner R, Endress G, Horrigan S;
PI Soppet DR, Weaver Z;
XX WPI; 2002-188264/24.
XX
PT Screening for anti-neoplastic agent involves exposing cells to a
PT chemical agent to be tested for anti-neoplastic activity, and
PT determining a change in expression of a gene of a signature gene set -
XX
XX Claim 1; SEQ ID 6897; 44pp; English.
XX
XX The present invention describes a method (M1) for screening for an
XX anti-neoplastic agent. The method involves exposing cells to a chemical
XX agent to be tested for anti-neoplastic activity, determining a change in
XX expression of at least one gene (I) of a signature gene set, where (I)
XX comprises a sequence (S) selected from 8447 sequences (given in ABL61664
XX to ABL70110), or is at least 95% identical to (S), where a change in
XX expression is indicative of anti-neoplastic activity. (I) has cytostatic
XX activity and can be used in gene therapy. M1 can be used for screening
XX an anti-neoplastic agent, and can be used for producing a product which
XX is the data collected with respect to the anti-neoplastic agent as a
XX result of M1, and the data is sufficient to convey the chemical
XX structure and/or properties of the agent. M1 can be used in the
XX treatment of cancer such as colon, breast, stomach, lung, thyroid,
XX oesophageal, ovarian, kidney, prostate or pancreatic cancer.
XX adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer,
XX infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine
XX carcinoma, papillary carcinoma and Wilms' tumour.
SQ Sequence 267156 BP; 76527 A; 56343 C; 55787 G; 78499 T; 0 other;

Query Match 5.9%; Score 58.6; DB 24; Length 267156;
Best Local Similarity 64.1%; Pred. No. 1.6e-06;
Matches 139; Conservative 0; Mismatches 69; Indels 9; Gaps 3;

QY 1 AGCAACCTGTAAGTTCGGCTGCAATCATAGATAGTAAGTGAAGCTTGTATGGGAG 60
III III III III III III III III III III III III III III III III III III
Db 43125 AGCATCCTGAAATATTTAGCTGCAACACATAGTTAAGAAAGCTTGCATGGGA- 43183

CC result of M1, and the data is sufficient to convey the chemical
CC structure and/or properties of the agent. M1 can be used in the
CC treatment of cancer such as colon, breast, stomach, lung, thyroid,
CC oesophageal, ovarian, kidney, prostate or pancreatic cancer,
CC adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer,
CC infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine
CC carcinoma, papillary carcinoma and Wilm's tumour.
XX

SQ Sequence 149480 BP; 39770 A; 34233 C; 35027 G; 40449 T; 1 other;
Query Match 4.9%; Score 48.8; DB 24; Length 149480;
Best Local Similarity 64.5%; Pred. No. 0.0016;
Matches 89; Conservative 0; Mismatches 47; Indels 2; Gaps 1;

Qy 1 AGCAACCTGTAAGTTCGGCTGCAATCATAGATAAGTAAGTGAAGCTTGTATGGGCAG 60
Db 55060 AGCAGCCAGTAAATCGAGCTGTAGACATAGTCAAGGGAGCTGGNAGCTTACACGGG--T 55117
Qy 61 GGATGGCTGCAGCTTCATGGATGAGAAATGTCCAGCTTGGGCTAGATACATCCAAACATGGG 120
Db 55118 GAATGCTGGCAGCTGTGCCCATAGGAAAAGGCCACCTGGTCTAGGTATGTTCAAAATGGC 55177
Qy 121 GGCCTCCACTCCTCTTGT 138
Db 55178 GGCTCCAGGCTCCCTTCT 55195

Search completed: December 24, 2002, 17:17:35
Job time : 558.952 secs

THIS PAGE BLANK (USPTO)

GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: December 24, 2002, 16:23:56 ; Search time 165.952 Seconds
(without alignments)
13583.723 Million cell updates/sec

Title: US-09-708-724A-3_COPY_10000_11000

Perfect score: 1001

Sequence: 1 caaaatttcagtaggaaga.....gcagcacataatgatcatg 1001

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 2185239 seqs, 1125999159 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 20000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

N_Geneseq_101002.*

- 1: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1980.DAT.*
- 2: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1981.DAT.*
- 3: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1982.DAT.*
- 4: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1983.DAT.*
- 5: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1984.DAT.*
- 6: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1985.DAT.*
- 7: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1986.DAT.*
- 8: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1987.DAT.*
- 9: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1988.DAT.*
- 10: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1989.DAT.*
- 11: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1990.DAT.*
- 12: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1991.DAT.*
- 13: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1992.DAT.*
- 14: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1993.DAT.*
- 15: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1994.DAT.*
- 16: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1995.DAT.*
- 17: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1996.DAT.*
- 18: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1997.DAT.*
- 19: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1998.DAT.*
- 20: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1999.DAT.*
- 21: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2000.DAT.*
- 22: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001A.DAT.*
- 23: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001B.DAT.*
- 24: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Length	ID	Description
1	82	8.2	119950	20 AAX90201 Human yes1 gene.
2	79	7.9	43069	21 AA236335 Genomic sequence o
3	77.2	7.7	368004	24 ABL57909 Human transporter
4	76	7.6	302250	24 ABL67703 Oesophagus cancer
5	75.8	7.6	17000	22 AAL40299 Caspase 6 antisens
6	75.2	7.5	1167	22 AAS00839 Human cDNA clone H
7	75.2	7.5	1231	22 AAH33190 Human colon cancer
8	73.2	7.3	199	21 AAC19473 Human secreted pro
9	73.2	7.3	6312	22 ABA14390 Human nervous syst

10	72.8	7.3	699	22 ABA07676 Human ovarian and
11	72.8	7.3	699	22 AAL02590 Human reproductive
12	72.6	7.3	2079	22 AAK94192 Human full-length
13	72.4	7.2	592	24 ABN60912 Human cancer relat
14	72.4	7.2	14968	21 AAF21343 Human low adenosin
15	72.4	7.2	14968	21 AAA35221 Human adenosine re
16	72.4	7.2	14968	22 AAD14464 Human IL-15 gene a
17	72.4	7.2	14968	22 AAD15838 Human interleukin
18	72.4	7.2	17844	21 AAA35223 Human adenosine re
19	72.4	7.2	17904	21 AAF21345 Human low adenosin
20	72.2	7.2	343	21 AAC29850 Human secreted pro
21	71.4	7.1	894	22 AAK64791 Human immune/haema
22	71.2	7.1	1885	22 AAH14899 Human cDNA sequenc
23	71.2	7.1	3208	22 AAI14896 Human cDNA sequenc
24	70.8	7.1	397	22 AAI89485 Human polynucleoti
25	70.8	7.1	465237	24 ABQ87681 Human oestrogen re
26	70.8	7.1	465237	24 ABA07681 Human oestrogen re
27	70.6	7.1	136284	24 ABK83575 Human cDNA differe
28	70.2	7.0	678	22 AAS46934 Human G protein-co
29	70.2	7.0	678	24 ABK81712 cDNA encoding nove
30	70.2	7.0	7687	22 AAL04658 Human reproductive
31	70.2	7.0	7687	23 ABL97565 Human testicular a
32	69.8	7.0	742	22 AAI95221 Human neuroblastom
33	69.4	6.9	560	24 ABL78544 Human ovarian canc
34	69.4	6.9	2934	22 ABA07330 Human pancreatic c
35	69.4	6.9	2934	22 AAK64817 Human immune/haema
36	69.4	6.9	2934	22 AAK90489 Human digestive sy
37	69.2	6.9	240825	22 AAF24497 Human PG-3 gene.
38	69	6.9	534	22 ABA07011 Human pancreatic c
39	69	6.9	534	22 AAK88496 Human digestive sy
40	69	6.9	3581	22 AAS39909 Genomic sequence #
41	69	6.9	3581	22 AAK72164 Human immune/haema
42	69	6.9	3581	22 AAK90353 Human digestive sy
43	68.2	6.8	168575	22 AAH21613 Human hypocrerin r
44	67.2	6.7	4582	22 AAK72026 Human immune/haema
45	67.2	6.7	8734	22 AAK72027 Human immune/haema

ALIGNMENTS

RESULT 1

AAX90201
ID AAX90201 standard; DNA; 119950 BP.

XX AAX90201;

AC AAX90201;

XX 23-SEP-1999 (first entry)

XX Human yes1 gene.

XX Human; yes1; diagnosis: neuropsychiatric disorder; BAD; schizophrenia;
XX bipolar affective disorder; attention deficit disorder;
XX schizoaffective disorder; unipolar affective disorder;
XX Huntington's disease; Parkinson's disease; manic-depression; ds.

OS Homo sapiens.

XX WO9935290-A1.

XX 15-JUL-1999.

XX 07-JAN-1999; 99WO-US00297.

XX 08-JAN-1998; 98US-0003944.

XX (MILL-) MILLENNIUM PHARM INC.

XX Chen H, Freimer NB;

XX WPI; 1999-444203/37.

XX P-PSDB; AAY24421.

```
PT Detection of a genetic mutation in the yes1 gene, useful for
XX diagnosis of a yes1 mediated neuropsychiatric disorder in a human
PS Claim 1; Fig 2; 110pp; English.
XX
CC The present invention describes a method for detecting a genetic
CC mutation in the yes1 gene for the diagnosis of a yes1 mediated
CC neuropsychiatric disorder in a human. The method comprises detecting the
CC presence or absence of a genetic mutation in the yes1 gene of the
CC subject, where the genetic mutation is a substitution, insertion or a
CC deletion and results in the production of a yes1 protein having an amino
CC acid sequence other than the wild-type yes1 amino acid sequence and the
CC presence of the genetic mutation identifies a subject that has or is at
CC risk for developing a yes1 mediated neuropsychiatric disorder. Compounds
CC that bind to the yes1 protein, alter the amount of the protein, or alter
CC the activity of the yes1 gene product, are useful for treating a yes1
CC mediated neuropsychiatric disorder. The disorders include Huntington's
CC disease, Parkinson's disease, and especially bipolar-affective disorder
CC (BAD) also known as bipolar mood disorder (BP) or manic-depressive
CC illness. The method distinguishes neuropsychiatric disorders from
CC neurological disorders, which enables more accurate evaluation and
CC prescription of medical treatment. The present sequence represents the
CC human yes1 cDNA sequence.
XX
SQ Sequence 119950 BP; 34471 A; 23730 C; 24660 G; 37033 T; 56 other;

Query Match      8.2%; Score 82; DB 20; Length 119950;
Best Local Similarity 65.1%; Pred. No. 5.8e-11;
Matches 138; Conservative 0; Mismatches 70; Indels. 4; Gaps 1;

QY 3 AAATTCAGTTAGGAGATAAGTCGAAGAGATCTATTGTTACTTGGTGACTACAGTTAT 62
   ||| || ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 79787 AAATTTAGATAGGAGGAGTAAGTTCAAGAGATCTATTCTAGAACACAGTTAATAACAAT 79846

QY 63 GTATCTGTCTTGACTATACAGTAGATTTCGAGTGTTCACACACAAAACATCATGG 122
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 79847 ATATCTGTCTTGAAAATTAACAAAATTTTAAGTCTTCTCACCAAAAATGA--- 79903

QY 123 GTATGTGAGTAATGCATGATCAAACTAGCTGGGTAAACCAATTCACAAATATGTGTGA 182
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 79904 -TATGTGATGCATGATGATGATTAATAGCTTACTTACCCATTCACATGTATATATA 79962

QY 183 TTTCAAACAGTACCATAAATGACAGACAATTT 214
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 79963 TTTCAAAGCATCATGTTGTATGCAATAAATAT 79994

RESULT 2
AAZ36335/c
ID AAZ36335 standard; DNA; 43069 BP.
XX
AC AAZ36335;
XX
DT 22-FEB-2000 (first entry)
XX
DE Genomic sequence of the 5-lipoxygenase activating protein (FLAP).
XX
KW Human; 5-lipoxygenase activating protein; FLAP; biallelic marker;
KW leukotriene pathway; genotype; haplotype; FLAP-related biallelic marker;
KW asthma; ss.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT misc_feature 1..7007
FT /tag= a
FT /note= "specifically claimed in claim 1"
FT misc_feature 1..7708
FT /tag= b
FT /note= "potential 5' regulatory region"
FT primer_bind 3851..3869
FT /tag= c
FT /note= "upstream amplification primer 10-517"
```

```
FT allele
FT primer_bind
FT /tag= d
FT 3927..3973
FT /tag= e
FT /note= "potential binding site for a probe"
FT complement (4171..4189)
FT /tag= f
FT /note= "downstream amplification primer 10-517"
FT 4120..4138
FT /tag= g
FT /note= "upstream amplification primer 10-518"
FT 4220..4266
FT /tag= h
FT /note= "potential binding site for a probe"
FT replace (4243, T)
FT /tag= i
FT 4289..4335
FT /tag= j
FT /note= "potential binding site for a probe"
FT replace (4312, G)
FT /tag= k
FT complement (4372..4390)
FT /tag= l
FT /note= "downstream amplification primer 10-518"
FT 4373..4391
FT /tag= m
FT /note= "upstream amplification primer 10-253"
FT 4467..4513
FT /tag= n
FT /note= "potential binding site for a probe"
FT replace (4490, G)
FT /tag= o
FT 4647..4693
FT /tag= p
FT /note= "potential binding site for a probe"
FT 4664..4710
FT /tag= q
FT /note= "potential binding site for a probe"
FT replace (4670, G)
FT /tag= r
FT replace (4687, C)
FT /tag= s
FT complement (4773..4792)
FT /tag= t
FT /note= "downstream amplification primer 10-253"
FT 4814..4833
FT /tag= u
FT /note= "upstream amplification primer 10-499"
FT 4945..4991
FT /tag= v
FT /note= "potential binding site for a probe"
FT 4956..4972
FT /tag= w
FT /note= "upstream amplification primer 10-500"
FT replace (4968, A)
FT /tag= x
FT complement (5026..5043)
FT /tag= y
FT /note= "downstream amplification primer 10-499"
FT 5117..5163
FT /tag= z
FT /note= "potential binding site for a probe"
FT replace (5140, T)
FT /tag= aa
FT 5190..5236
FT /tag= ab
FT /note= "potential binding site for a probe"
FT replace (5213, G)
FT /tag= ac
FT 5341..5387
FT /tag= ad
FT /note= "potential binding site for a probe"
FT replace (5364, A)
```

```
FT primer_bind /tag= ae complement (5405..5422)
FT /tag= af
FT /note= "downstream amplification primer 10-500"
FT primer_bind 5524..5542
FT /tag= ag
FT /note= "upstream amplification primer 10-522"
FT primer_bind 5571..5617
FT /tag= ah
FT /note= "potential binding site for a probe"
FT allele replace (5594, A)
FT /tag= ai
FT primer_bind complement (5978..5996)
FT /tag= aj
FT /note= "downstream amplification primer 10-522"
FT primer_bind 6218..6235
FT /tag= ak
FT /note= "upstream amplification primer 10-503"
FT primer_bind 6347..6393
FT /tag= al
FT /note= "potential binding site for a probe"
FT allele replace (6370, G)
FT /tag= am
FT primer_bind 6522..6539
FT /tag= an
FT /note= "upstream amplification primer 10-504"
FT primer_bind complement (6652..6672)
FT /tag= ao
FT /note= "downstream amplification primer 10-503"
FT primer_bind 6670..6716
FT /tag= ap
FT /note= "potential binding site for a probe"
FT allele replace (6693, T)
FT /tag= aq
FT primer_bind 6740..6786
FT /tag= ar
FT /note= "potential binding site for a probe"
FT allele replace (6763, A)
FT /tag= as
FT primer_bind complement (6772..6790)
FT /tag= at
FT /note= "downstream amplification primer 10-504"
FT primer_bind 7120..7137
FT /tag= au
FT /note= "upstream amplification primer 10-204"
FT primer_bind 7422..7468
FT /tag= av
FT /note= "potential binding site for a probe"
FT allele replace (7445, A)
FT /tag= aw
FT primer_bind /note= "claim 4"
FT 7513..7531
FT /tag= ax
FT /note= "upstream amplification prior 10-32"
FT primer_bind complement (7557..7574)
FT /tag= ay
FT /note= "downstream amplification primer 10-204"
FT misc_feature 7612..7637
FT /tag= az
FT /note= "specifically claimed in claim 3"
FT exon 7709..7852
FT /tag= ba
FT /number= 1
FT CDS 7783..36250
FT /tag= bb
FT /product= FLAP
FT /note= "contains introns"
FT primer_bind 7847..7893
FT /tag= bc
FT /note= "potential binding site for a probe"
FT intron 7853..16235
FT /tag= bd
FT /number= 1
```

```
FT allele replace (7870, A)
FT /tag= be
FT /note= "claim 4"
FT primer_bind complement (7914..7933)
FT /tag= bf
FT /note= "downstream amplification primer 10-32"
FT misc_feature 8117..15994
FT /tag= bg
FT /note= "specifically claimed in claim 1"
FT primer_bind 16114..16132
FT /tag= bh
FT /note= "upstream amplification primer 10-33"
FT exon 16236..16335
FT /tag= bi
FT /number= 2
FT primer_bind 16265..16311
FT /tag= bj
FT /note= "potential binding site for a probe"
FT allele replace (16288, T)
FT /tag= bk
FT /note= "claim 4"
FT primer_bind 16324..16370
FT /tag= bl
FT /note= "potential binding site for a probe"
FT intron 16336..24226
FT /tag= bm
FT /number= 2
FT allele replace (16347, A)
FT /tag= bn
FT primer_bind 16360..16406

Query Match 7.9%; Score 79; DB 21; Length 43069;
Best Local Similarity 64.68; Pred. No. 2.4e-10;
Matches 135; Conservative 0; Mismatches 70; Indels 4; Gaps 1;

Qy 223 TTACAATCAAAAAGTTTAAATGAGGACCTTAGGGTGGTCTTAATCAATCTTAAGTG 282
Dy 34050 TTAAGAGATAATTAAGGTAAATCACTCATATGGTGGTCCCTCATCCAGTATGACTG 33991
Qy 283 ATGCTCCATGAAGAGGAATAAGGATACAAATGTGCACACAGAGAGAAATGGCCACAT 342
Dy 33990 GGGCTTTATAAGAGGGGAGATTTGGACACAGACATGCTCAGTGAGG---TGATGATGT 33935
Qy 343 GAGACACAATGAGAATGTGCTACTTACAGCCTAGGAGAGAGCGCTCCGAGAGAAACAC 402
Dy 33934 GAAGACACAGGTAGAGGTGGCACTCTATAGCCAAAGGAGAGAGGCTGCAGGAGAAACCG 33875
Qy 403 ACCCTACCCACACCTTGATGTTGGACTTC 431
Dy 33874 ACCCTGCCACACTTTGATCTTGGACTTC 33846

RESULT 3
ABL57909/c
ID .ABL57909 standard; DNA; 368004 BP.
XX AC ABL57909;
XX AC ABL57909;
DT 05-JUL-2002 (first entry)
XX DE Human transporter protein gene.
XX DE Human; transporter protein; gene; gene therapy; glutamate receptor 4; SNP;
XX KW brain; foetal brain; chromosome 11; single nucleotide polymorphism; ds.
XX OS Homo sapiens.
XX FH Key
XX FH CDS 2001..366005
XX FT /tag= a
XX FT /product= "Human transporter protein"
XX FT exon 2001..2088
XX FT /tag= b
```

```
FT /number= 1 /number= 1 variation replace(27764,C)
FT 3279...3437 /*tag= aa /*tag= aa
FT /*tag= c /*standard_name= "Single nucleotide polymorphism"
FT /number= 2 replace(27766,C)
FT 144533...144772 /*tag= ab
FT /*tag= d /*standard_name= "Single nucleotide polymorphism"
FT /number= 3 replace(28939,A)
FT 253472...253656 /*tag= ac
FT /*tag= e /*standard_name= "Single nucleotide polymorphism"
FT /number= 4 replace(29021,C)
FT 278947...279000 /*tag= ad
FT /*tag= f /*standard_name= "Single nucleotide polymorphism"
FT /number= 5 replace(29908,A)
FT 289708...289866 /*tag= ae
FT /*tag= g /*standard_name= "Single nucleotide polymorphism"
FT /number= 6 replace(30398,T)
FT 295254...295421 /*tag= af
FT /*tag= h /*standard_name= "Single nucleotide polymorphism"
FT /number= 7 replace(30440,T)
FT 296637...296741 /*tag= ag
FT /*tag= i /*standard_name= "Single nucleotide polymorphism"
FT /number= 8 replace(31718,G)
FT 301876...301986 /*tag= ah
FT /*tag= j /*standard_name= "Single nucleotide polymorphism"
FT /number= 9 replace(35206,C)
FT 310162...310368 /*tag= ai
FT /*tag= k /*standard_name= "Single nucleotide polymorphism"
FT /number= 10 replace(35304,A)
FT 315851...316221 /*tag= aj
FT /*tag= l /*standard_name= "Single nucleotide polymorphism"
FT /number= 11 replace(36362,A)
FT 318193...318391 /*tag= ak
FT /*tag= m /*standard_name= "Single nucleotide polymorphism"
FT /number= 12 replace(36436,A)
FT 325176...325423 /*tag= al
FT /*tag= n /*standard_name= "Single nucleotide polymorphism"
FT /number= 13 replace(43063,C)
FT 357392...357532 /*tag= am
FT /*tag= o /*standard_name= "Single nucleotide polymorphism"
FT /number= 14 replace(43202,C)
FT 363364...363478 /*tag= an
FT /*tag= p /*standard_name= "Single nucleotide polymorphism"
FT /number= 15 replace(44576,G)
FT 365760...366002 /*tag= ao
FT /*tag= q /*standard_name= "Single nucleotide polymorphism"
FT /number= 16 replace(44622,G)
FT replace(577,C) /*tag= ap
FT /*tag= r /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(45378,T)
FT replace(1895,G) /*tag= aq
FT /*tag= s /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(45685,T)
FT replace(2765,A) /*tag= ar
FT /*tag= t /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(45998,T)
FT replace(10364,G) /*tag= as
FT /*tag= u /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(46173,C)
FT replace(11079,G) /*tag= at
FT /*tag= v /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(47636,A)
FT replace(11514,G) /*tag= au
FT /*tag= w /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(48011,G)
FT replace(20503,G) /*tag= av
FT /*tag= x /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(48012,C)
FT replace(20505,G) /*tag= aw
FT /*tag= y /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(48019,A)
FT replace(23307,A) /*tag= ax
FT /*tag= z /*standard_name= "Single nucleotide polymorphism"
FT /standard_name= "Single nucleotide polymorphism" replace(50175,G)
```



```
FT FT variation /*tag= ay /standard_name= "Single nucleotide polymorphism"
FT FT replace(50919,T)
FT FT /*tag= az /standard_name= "Single nucleotide polymorphism"
FT FT /*tag= ba replace(51730,T)
FT FT /*tag= bb /standard_name= "Single nucleotide polymorphism"
FT FT replace(53975,G)
FT FT /*tag= bc /standard_name= "Single nucleotide polymorphism"
FT FT replace(54377,C)
FT FT /*tag= bd /standard_name= "Single nucleotide polymorphism"
FT FT replace(54388,G)
FT FT /*tag= be /standard_name= "Single nucleotide polymorphism"
FT FT replace(61997,G)
FT FT /*tag= bf /standard_name= "Single nucleotide polymorphism"
FT FT replace(62272,C)
FT FT /*tag= bg /standard_name= "Single nucleotide polymorphism"
FT FT replace(63212,G)
FT FT /*tag= bh /standard_name= "Single nucleotide polymorphism"
FT FT replace(65792,C)
FT FT /*tag= bi /standard_name= "Single nucleotide polymorphism"
FT FT replace(69135,G)
FT FT /*tag= bj /standard_name= "Single nucleotide polymorphism"
FT FT /*tag= bk /standard_name= "Single nucleotide polymorphism"

Query Match 7.7%; Score 77.2; DB 24; Length 368004;
Best Local Similarity 68.5%; Pred. No. 1.7e-09;
Matches 137; Conservative 0; Mismatches 59; Indels 5; Gaps 2;

QY 231 RAAAAAGTTTAAATAGGACCTTAGGGTGGTCCCTTAATCCAAATCAAGTATGCTCC 290
Db 33099 AGAATCGAGTTAAATAGGTCATTAGGGTGGCCCTTAATGCAATATGACTGGTCTTC 33040

QY 291 ATGAAGAGGAATAGATACAAATGTGCACAGAGAGAAATGCCACATGAGGACAC 350
Db 33039 ATAAAGAGGAATAGGACACAGA----CATGCAGAGAGGGAAGACCATATTAAGACAC 32984

QY 351 AATGAGATGTGGCTACTTACAGCCTAGGAGAGGCGCTCCGAGAAACACACCCCTACC 410
Db 32983 AGGGAGATAATGGC-ATCTACAAGCAAGGAGAAAGGCGCTCAGAAACCAACCTTTGCT 32925

QY 411 CACACCTTGATCTTGACTT 430
Db 32924 GCCTCCTTGATCTCAGACTT 32905

RESULT 4
ABL67703
ID ABL67703 standard; DNA; 302250 BP.
XX AC ABL67703;
XX XX
XX 15-MAY-2002 (first entry)
XX DE Oesophagus cancer related gene sequence SEQ ID NO:6040.
XX KW Human; cancer; colon; breast; ovary; oesophagus; kidney; thyroid;
XX KW stomach; lung; prostate; pancreas; carcinoma; antitumour; cancerous;
XX KW cystostatic; gene therapy; antineoplastic; Wilms' tumour; adenocarcinoma;
XX KW gene; ds.
XX OS Homo sapiens.
```

```
XX WO200194629-A2.
XX 13-DEC-2001.
XX 30-MAY-2001; 2001WO-US10838.
XX 05-JUN-2000; 2000US-209473P.
XX 05-JUN-2000; 2000US-209531P.
XX 18-SEP-2000; 2000US-233133P.
XX 18-SEP-2000; 2000US-233617P.
XX 20-SEP-2000; 2000US-234009P.
XX 20-SEP-2000; 2000US-234034P.
XX 22-SEP-2000; 2000US-234509P.
XX 22-SEP-2000; 2000US-234567P.
XX 25-SEP-2000; 2000US-234923P.
XX 25-SEP-2000; 2000US-234924P.
XX 25-SEP-2000; 2000US-235077P.
XX 25-SEP-2000; 2000US-235082P.
XX 25-SEP-2000; 2000US-235134P.
XX 25-SEP-2000; 2000US-235280P.
XX 26-SEP-2000; 2000US-235637P.
XX 26-SEP-2000; 2000US-235638P.
XX 27-SEP-2000; 2000US-235711P.
XX 27-SEP-2000; 2000US-235720P.
XX 27-SEP-2000; 2000US-235840P.
XX 27-SEP-2000; 2000US-235863P.
XX 28-SEP-2000; 2000US-236028P.
XX 28-SEP-2000; 2000US-236032P.
XX 28-SEP-2000; 2000US-236033P.
XX 28-SEP-2000; 2000US-236034P.
XX 28-SEP-2000; 2000US-236109P.
XX 28-SEP-2000; 2000US-236111P.
XX 29-SEP-2000; 2000US-236842P.
XX 29-SEP-2000; 2000US-236891P.
XX 02-OCT-2000; 2000US-237172P.
XX 02-OCT-2000; 2000US-237173P.
XX 02-OCT-2000; 2000US-237278P.
XX 02-OCT-2000; 2000US-237294P.
XX 02-OCT-2000; 2000US-237295P.
XX 02-OCT-2000; 2000US-237316P.
XX 03-OCT-2000; 2000US-237425P.
XX 03-OCT-2000; 2000US-237598P.
XX 03-OCT-2000; 2000US-237604P.
XX 03-OCT-2000; 2000US-237606P.
XX 03-OCT-2000; 2000US-237608P.
XX 01-NOV-2000; 2000US-244867P.
XX 01-NOV-2000; 2000US-245084P.
XX (AVAL-) AVALON PHARM.
XX Young PE, Augustus M, Carter KC, Ebner R, Endress G, Horrigan S;
PI Soppet DR, Weaver Z;
XX WPI; 2002-188264/24.
XX Screening for anti-neoplastic agent involves exposing cells to a
PT chemical agent to be tested for anti-neoplastic activity, and
PT determining a change in expression of a gene of a signature gene set
XX Claim 1; SEQ ID 6040; 44pp; English.
XX The present invention describes a method (M1) for screening for an
CC anti-neoplastic agent. The method involves exposing cells to a chemical
CC agent to be tested for anti-neoplastic activity, determining a change in
CC expression of at least one gene (I) of a signature gene set, where (I)
CC comprises a sequence (S) selected from 8447 sequences (given in ABL61664
CC to ABL70110), or is at least 95% identical to (S), where a change in
CC expression is indicative of anti-neoplastic activity. (I) has cytostatic
CC activity and can be used in gene therapy. M1 can be used for screening
CC an anti-neoplastic agent, and can be used for producing a product which
CC is the data collected with respect to the anti-neoplastic agent as a
```

CC result of M1, and the data is sufficient to convey the chemical
CC structure and/or properties of the agent. M1 can be used in the
CC treatment of cancer such as colon, breast, stomach, lung, thyroid,
CC oesophageal, ovarian, kidney, prostate or pancreatic cancer,
CC adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer,
CC infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine
CC carcinoma, papillary carcinoma and Wilms' tumour.
XX

SQ Sequence 302250 BP; 76116 A; 72066 C; 71554 G; 82514 T; 0 other;
Query Match 7.6%; Score 76; DB 24; Length 302250;
Best Local Similarity 65.6%; Pred. No. 3.3e-09;
Matches 143; Conservative 0; Mismatches 70; Indels 5; Gaps 2;

QY 215 TGTGTCAGTTCAATCAAAAAGTTTAAATGAGGACCTTAGGTGGTCTTAATCCAA 274
DB 113304 TGGGGCTTTAATAGTAAATAGATAGATGAGCGCTTAGGTGGTCTTAATCCAA 113363
QY 275 TCTAGTGTAGT-CTCCATGAAGAGGAAATAGGATACAAATGTCACACAGAGAGAA 333
DB 113364 TATGACTGGTGTCCATGTTAAAAAGGAGATTTCAGACACAGACTTGTGCAGAGGGAGAA- 113422
QY 334 TGGCCACATGAGACACATGAGATGTGGCTACTTACAAGCCTAGGAGAGGCGCTCCG 393
DB 113423 ---CCATGTGGGACGCGAGGAGAGTGGGCCATCTACAAGCCAGGACAGAGGCGCTCAG 113479
QY 394 AGAAACACACACCTACCCACACCTTGTGATGTGGACTTC 431
DB 113480 AATGAACCAACCTGCCACACCTTGGTCTCCAACTTC 113517

RESULT 5
AAL40299/c
ID AAL40299 standard; DNA; 17000 BP.

XX AAL40299;
XX
DT 19-SEP-2002 (first entry)
XX
DE Caspase 6 antisense inhibition related nucleic acid SEQ ID No 18.
KW Muscular; cytostatic; nootropic; neuroprotective; ophthalmological;
KW antilipemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism;
KW ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder;
KW haematopoietic disorder; cancer; neurological; Alzheimer's disease;
KW apoptotic; human; gene; ds.
XX Homo sapiens.
XX WO200229066-A1.
XX
PD 11-APR-2002.
XX
PF 03-OCT-2001; 2001WO-US30871.
XX
PR 04-OCT-2000; 2000US-0679299.
XX
PA (ISIS-) ISIS PHARM INC.
XX
PI Brown-driver VL, Zhang H, Watt AT;
XX
XX WPI; 2002-471315/50.
DR
PT An antisense oligonucleotide of 8 to 50 nucleotides in length that
XX inhibits caspase 6, is useful for treating Rieger's syndrome -
XX
XX Example 15; Page 104-112; 141pp; English.

XX The invention relates to an antisense oligonucleotide compound of 8 to 50
CC nucleotides in length that is targeted to a nucleic acid molecule
CC encoding caspase 6, where the oligonucleotide specifically hybridises
CC with and inhibits the expression of caspase 6. The oligonucleotide of the
CC invention specifically hybridises to and inhibits expression of caspase 6

CC in cells or tissues. The oligonucleotides can be administered
CC therapeutically or prophylactically to treat an animal having a disease
CC or condition associated with caspase 6, such as Rieger's syndrome or
CC ataxia telangiectasia, hyperproliferative disorder, a haematopoietic
CC disorder, a bone metabolism or cholesterol disorder, various types of
CC cancer, neurological conditions such as Alzheimer's disease and other de-
CC regulated apoptotic pathological conditions. This polynucleotide sequence
CC represents a human caspase 6 nucleic acid relating to the invention.
XX

SQ Sequence 17000 BP; 4758 A; 3480 C; 3987 G; 4775 T; 0 other;
Query Match 7.6%; Score 75.8; DB 24; Length 17000;
Best Local Similarity 68.3%; Pred. No. 1.2e-09;
Matches 136; Conservative 0; Mismatches 57; Indels 6; Gaps 2;

QY 242 AAAATGAGGACCTTAGGTGGTCTTAATCCAATTAAGTGTATGCTCCATGAAGAGGA 301
DB 12724 AAAATGAGGTCTAGCGGTGAACCAATCAACATGACTAGTGTCTTCAT--AGAAGA 12667
QY 302 AATAAGGATACAAATGTCACACAGAGAGAAATGGCCACATGAGGACACAATGAGAATGT 361
DB 12666 AATTGGGACACAGACATGCAC----AGAAAGAACCATATGAAGGCACAGGACAAGAT 12611
QY 362 GCGTACTTACAGCCTAGGAGAGAGCGCTCCGAGAAACACACACCTACCCACACCTTGTAT 421
DB 12610 GGCCACCTACAAAGCCAAAAGGGGCGCTCAGAAGAAACCAATCTCCCCACACGCTTCAT 12551
QY 422 GTTGGACTTCATCCGTAG 440
DB 12550 CTGAGACTTCAGCTCCAG 12532

RESULT 6
AAS00839/c
ID AAS00839 standard; cDNA; 1167 BP.

XX AAS00839;
XX
DT 04-JUL-2001 (first entry)
XX
DE Human cDNA clone HWLEH32 encoding cancer related protein 13.
KW Human; cancer related protein; HWLEH32; food additive;
KW preservative; immunogen; antibody; bone cancer; adrenal cancer;
KW bone marrow cancer; breast cancer; gastrointestinal cancer;
KW liver cancer; lung cancer; urogenital cancer; immune disorder;
KW Addison's disease; allergy; autoimmune haemolytic anaemia;
KW autoimmune thyroiditis; diabetes mellitus; Crohn's disease;
KW multiple sclerosis; rheumatoid arthritis; ulcerative colitis;
KW acquired immunodeficiency syndrome; AIDS; cardiovascular disorder;
KW myocardial ischaemia; wound healing; neurological disorder;
KW Parkinson's disease; Alzheimer's disease; cerebral anoxia; epilepsy;
KW viral infection; bacterial infection; fungal infection;
KW parasitic infection; agonist; antagonist; ss.

XX Homo sapiens.
XX
XX Key Location/Qualifiers
XX CDS 592..741
XX /tag= a
XX /product= "Cancer related protein 13"
XX sig_peptide 592..633
XX /tag= b
XX mat_peptide 634..738
XX /tag= c
XX /label= Mature_Cancer_related_protein_13

XX WO200118014-A1.
XX
XX 15-MAR-2001.
XX
XX 30-AUG-2000; 2000WO-US23794.
XX

XX 06-SEP-2000.
XX 21-FEB-2000; 2000EP-0200610.
XX 26-FEB-1999; 99US-0122487.
XX (GEST) GENSET.
XX Dumas Milne Edwards J, Duclert A, Giordano J;
XX WPI; 2000-500381/45.
XX New nucleic acid that is a 5' expressed sequence tag (5' EST) for
PT obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for
PT diagnostic, forensic, gene therapy and chromosome mapping procedures -
XX Claim 1; SEQ ID 23548; 7lpp + CD-ROM; English.
XX The present sequence is one of a large number of 5' ESTs derived from
CC mRNAs encoding secreted proteins. No ORF has yet been conclusively
CC identified within the present sequence. The 5' ESTs were prepared from
CC total human RNAs or polyA+ RNAs derived from 30 different tissues. EST
CC sequences usually correspond mainly to the 3' untranslated region (UTR)
CC of the mRNA because they are often obtained from oligo-dT primed cDNA
CC libraries. Such ESTs are not well suited for isolating cDNA sequences
CC derived from the 5' ends of mRNAs and even in those cases where longer
CC cDNA sequences have been obtained, the full 5' UTR is rarely included.
CC 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be
CC used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used
CC in diagnostic, forensic, gene therapy and chromosome mapping procedures.
CC They are used to obtain upstream regulatory sequences and to design
CC expression and secretion vectors.
XX Sequence 199 BP; 74 A; 34 C; 53 G; 38 T; 0 other;
Query Match 7.3%; Score 73.2; DB 21; Length 199;
Best Local Similarity 65.1%; Pred. No. 9.8e-10;
Matches 108; Conservative 0; Mismatches 58; Indels 0; Gaps 0;
QY 241 TAAATGAGGACCTAGGTGGTCTATCCCAATCTAGTGTCTCCATGAAAGAGG 300
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
30 TAAATGAGGTTCTATGGATGGGCCCTAATCCCAATGACTGTCTTAAAGAGGAGG 89
QY 301 AAATAGGATACAAATGTGCACACAGACAGAGAAATGCCACATGAGGACACAAATGAGAATG 360
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
90 AAATAGGACACAGACACACAAAGACTGAGGACGCTATGTGGGACAGACAGAAATG 149
QY 361 TGGCTACTTTACAGGCTAGGAGAGGCGCTCCGAGAAACACACCC 406
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
150 TGGCCATTGCAACCAAGGAGAGGCGCTCTGAAGAAACCAAC 195
RESULT 9
ID ABA14390
XX ABA14390 standard; DNA; 6312 BP.
XX AC ABA14390;
XX DT 23-JAN-2002 (first entry)
XX Human nervous system related polynucleotide SEQ ID NO 6721.
DE Human
XX Human
KW Human; nontropic; neuroprotective; cytostatic; dermatological; virucide;
KW immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary;
KW antiparkinsonian; antiskinking; antianaemic; antiarthritic; cancer;
KW antirheumatic; hepatotropic; cerebroprotective; antiinflammatory;
KW antiallergic; antidiabetic; antitumor; anticonvulsant; antifungal;
KW antiparasitic; cardiant; immune disorder; cardiovascular disorder;
KW neurological disease; infection; nephrotropic; gene therapy; vaccine; ds.
XX Homo sapiens.
OS
XX

PN WO200159063-A2.
XX 16-AUG-2001.
XX 17-JAN-2001; 2001WO-US01334.
XX 31-JAN-2000; 2000US-0179065.
PR 04-FEB-2000; 2000US-0180628.
PR 24-FEB-2000; 2000US-0184664.
PR 02-MAR-2000; 2000US-0186350.
PR 16-MAR-2000; 2000US-0189874.
PR 17-MAR-2000; 2000US-0190076.
PR 18-APR-2000; 2000US-0198123.
PR 19-MAY-2000; 2000US-0205515.
PR 07-JUN-2000; 2000US-0209467.
PR 28-JUN-2000; 2000US-0214886.
PR 30-JUN-2000; 2000US-0215135.
PR 07-JUL-2000; 2000US-0216647.
PR 11-JUL-2000; 2000US-0216880.
PR 11-JUL-2000; 2000US-0217496.
PR 14-JUL-2000; 2000US-0218290.
PR 26-JUL-2000; 2000US-0220964.
PR 26-JUL-2000; 2000US-0220964.
PR 14-AUG-2000; 2000US-0224519.
PR 14-AUG-2000; 2000US-0225213.
PR 14-AUG-2000; 2000US-0225214.
PR 14-AUG-2000; 2000US-0225266.
PR 14-AUG-2000; 2000US-0225267.
PR 14-AUG-2000; 2000US-0225268.
PR 14-AUG-2000; 2000US-0225270.
PR 14-AUG-2000; 2000US-0225447.
PR 14-AUG-2000; 2000US-0225757.
PR 14-AUG-2000; 2000US-0225758.
PR 14-AUG-2000; 2000US-0225759.
PR 18-AUG-2000; 2000US-0226279.
PR 22-AUG-2000; 2000US-0226681.
PR 22-AUG-2000; 2000US-0226688.
PR 22-AUG-2000; 2000US-0227182.
PR 23-AUG-2000; 2000US-0227009.
PR 30-AUG-2000; 2000US-0228924.
PR 01-SEP-2000; 2000US-0229287.
PR 01-SEP-2000; 2000US-0229343.
PR 01-SEP-2000; 2000US-0229344.
PR 01-SEP-2000; 2000US-0229345.
PR 05-SEP-2000; 2000US-0229509.
PR 05-SEP-2000; 2000US-0229513.
PR 06-SEP-2000; 2000US-0230437.
PR 08-SEP-2000; 2000US-0230438.
PR 08-SEP-2000; 2000US-0231242.
PR 08-SEP-2000; 2000US-0231243.
PR 08-SEP-2000; 2000US-0231244.
PR 08-SEP-2000; 2000US-0231413.
PR 08-SEP-2000; 2000US-0231414.
PR 08-SEP-2000; 2000US-0232080.
PR 08-SEP-2000; 2000US-0232081.
PR 12-SEP-2000; 2000US-0231968.
PR 14-SEP-2000; 2000US-0232397.
PR 14-SEP-2000; 2000US-0232398.
PR 14-SEP-2000; 2000US-0232399.
PR 14-SEP-2000; 2000US-0232400.
PR 14-SEP-2000; 2000US-0232401.
PR 14-SEP-2000; 2000US-0233063.
PR 14-SEP-2000; 2000US-0233064.
PR 14-SEP-2000; 2000US-0233065.
PR 21-SEP-2000; 2000US-0234223.
PR 21-SEP-2000; 2000US-0234274.
PR 25-SEP-2000; 2000US-0234997.
PR 25-SEP-2000; 2000US-0234998.
PR 26-SEP-2000; 2000US-0235484.
PR 27-SEP-2000; 2000US-0235834.
PR 27-SEP-2000; 2000US-0235836.

PR 29-SEP-2000; 2000US-0236327.
PR 29-SEP-2000; 2000US-0236367.
PR 29-SEP-2000; 2000US-0236368.
PR 29-SEP-2000; 2000US-0236369.
PR 29-SEP-2000; 2000US-0236370.
PR 02-OCT-2000; 2000US-0236802.
PR 02-OCT-2000; 2000US-0237037.
PR 02-OCT-2000; 2000US-0237038.
PR 02-OCT-2000; 2000US-0237039.
PR 02-OCT-2000; 2000US-0237040.
PR 13-OCT-2000; 2000US-0239935.
PR 13-OCT-2000; 2000US-0239937.
PR 20-OCT-2000; 2000US-0240960.
PR 20-OCT-2000; 2000US-0241785.
PR 20-OCT-2000; 2000US-0241786.
PR 20-OCT-2000; 2000US-0241787.
PR 20-OCT-2000; 2000US-0241808.
PR 20-OCT-2000; 2000US-0241809.
PR 20-OCT-2000; 2000US-0241826.
PR 20-OCT-2000; 2000US-0242221.
PR 01-NOV-2000; 2000US-0244617.
PR 08-NOV-2000; 2000US-0246474.
PR 08-NOV-2000; 2000US-0246475.
PR 08-NOV-2000; 2000US-0246476.
PR 08-NOV-2000; 2000US-0246477.
PR 08-NOV-2000; 2000US-0246478.
PR 08-NOV-2000; 2000US-0246523.
PR 08-NOV-2000; 2000US-0246524.
PR 08-NOV-2000; 2000US-0246525.
PR 08-NOV-2000; 2000US-0246526.
PR 08-NOV-2000; 2000US-0246527.
PR 08-NOV-2000; 2000US-0246528.
PR 08-NOV-2000; 2000US-0246532.
PR 08-NOV-2000; 2000US-0246609.
PR 08-NOV-2000; 2000US-0246610.
PR 08-NOV-2000; 2000US-0246611.
PR 08-NOV-2000; 2000US-0246613.
PR 17-NOV-2000; 2000US-0249207.
PR 17-NOV-2000; 2000US-0249208.
PR 17-NOV-2000; 2000US-0249209.
PR 17-NOV-2000; 2000US-0249210.
PR 17-NOV-2000; 2000US-0249211.
PR 17-NOV-2000; 2000US-0249212.
PR 17-NOV-2000; 2000US-0249213.
PR 17-NOV-2000; 2000US-0249214.
PR 17-NOV-2000; 2000US-0249215.
PR 17-NOV-2000; 2000US-0249216.
PR 17-NOV-2000; 2000US-0249217.
PR 17-NOV-2000; 2000US-0249218.
PR 17-NOV-2000; 2000US-0249244.
PR 17-NOV-2000; 2000US-0249245.
PR 17-NOV-2000; 2000US-0249264.
PR 17-NOV-2000; 2000US-0249265.
PR 17-NOV-2000; 2000US-0249267.
PR 17-NOV-2000; 2000US-0249299.
PR 17-NOV-2000; 2000US-0249300.
PR 01-DEC-2000; 2000US-0250391.
PR 01-DEC-2000; 2000US-0251160.
PR 05-DEC-2000; 2000US-0251030.
PR 05-DEC-2000; 2000US-0251988.
PR 05-DEC-2000; 2000US-0256719.
PR 06-DEC-2000; 2000US-0251479.
PR 08-DEC-2000; 2000US-0251856.
PR 08-DEC-2000; 2000US-0251868.
PR 08-DEC-2000; 2000US-0251869.
PR 08-DEC-2000; 2000US-0251989.
PR 08-DEC-2000; 2000US-0251990.
PR 11-DEC-2000; 2000US-0254097.
PR 05-JAN-2001; 2001US-0259678.
PR (HUMA-) HUMAN GENOME SCI INC.
PA Rosen CA, Barash SC, Ruben SM;
XX
PI

XX WPI; 2001-541565/60.
DR Nucleic acids encoding 3224 human nervous system antigen polypeptides,
XX useful for preventing, diagnosing and/or treating nervous system
PT cancers and metastases
PT
XX
PS Disclosure; SEQ ID NO 6721; 1701pp + Sequence Listing; English.
XX
XX The invention relates to novel genes (ABAI1004-ABA21534) and proteins
CC (ABBI4678-ABBI18001) useful for preventing, treating or ameliorating
CC medical conditions e.g. by protein or gene therapy. The genes are
CC isolated from a range of human tissues disclosed in the specification.
CC The nucleic acids, proteins, antibodies and (ant)agonists are useful
CC in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast
CC and ovarian cancer and other cancers of the adrenal gland, bone, bone
CC marrow, breast, gastrointestinal tract, liver, lung, or urogenital;
CC (b) immune disorders e.g. Addison's disease, allergies, autoimmune
CC haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's
CC disease, multiple sclerosis, rheumatoid arthritis and ulcerative
CC colitis; (c) cardiovascular disorders such as myocardial ischaemias;
CC (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and
CC epilepsy; and (f) infectious diseases such as viral, bacterial, fungal
CC and parasitic infections.
CC Note: The sequence data for this patent did not form part of the
CC printed specification, but was obtained in electronic format directly
CC from WIPO at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 6312 BP; 1750 A; 1411 C; 1347 G; 1804 T; 0 other;
Query Match 7.3%; Score 73.2; DB 22; Length 6312;
Best Local Similarity 65.4%; Pred. No. 3.9e-09;
Matches 140; Conservative 0; Mismatches 68; Indels 6; Gaps 2;
QY 218 GTCAGTTACAAATCAAAAAGTTTAAATAGAGACCTTAGGGTGGTCTCTTAATCAATCT 277
DB 6061 GGCCTTTAAAGAGAGTAAATTAATTTATACGGGGTGTGGTGAACCTTAATCAATAT 6120
QY 278 AAGTGATGTCCTCCAGAGAGAAATGAAGATACAATGTGCACACAGAGAAATGCG 337
DB 6121 GATTG--GTATCCTTGTAGGAGAGATTAGGACACAAA---CAACAGACAGAGGTGAC 6174
QY 338 CACATGAGGACACAATGAGAATGTGGCTACTTACAAGCCTAGGAGAGCGCTCCGAGAA 397
DB 6175 CACAGGAGGACACAGTGAGAAGCTGGCCATCTGCAAGCAAGGAGAGAGACCTCAGGAGA 6234
QY 398 AACACACCTTACCCACACCTTGATGTTGGACTTC 431
DB 6235 AACCAACCTGCCAGCACCTTAATCTTAGACTTC 6268
RESULT 10
ABA07676
ID ABA07676 standard; cDNA; 699 BP.
XX
AC ABA07676;
XX
DT 11-JAN-2002 (first entry)
XX
XX Human ovarian and breast cancer associated polynucleotide SEQ ID NO 233.
XX
XX Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral;
KW antiallergic; hepatotropic; antidiabetic; antinflammatory; antitumor;
KW vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic;
KW cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
KW neurological disease; infection; human; secreted protein; ss.
XX
OS Homo sapiens.
XX
XX WO200155325-A2.
PN
XX
PD 02-AUG-2001.
XX

PF 17-JAN-2001; 2001WO-US01345.
XX 31-JAN-2000; 2000US-0179065.
PR 04-FEB-2000; 2000US-0180628.
PR 24-FEB-2000; 2000US-0184664.
PR 02-MAR-2000; 2000US-0186350.
PR 16-MAR-2000; 2000US-0189874.
PR 17-MAR-2000; 2000US-0190076.
PR 18-APR-2000; 2000US-0198123.
PR 19-MAY-2000; 2000US-0205515.
PR 07-JUN-2000; 2000US-0209467.
PR 28-JUN-2000; 2000US-0214886.
PR 30-JUN-2000; 2000US-0215135.
PR 07-JUL-2000; 2000US-0216647.
PR 07-JUL-2000; 2000US-0216880.
PR 11-JUL-2000; 2000US-0217487.
PR 11-JUL-2000; 2000US-0217496.
PR 14-JUL-2000; 2000US-0218290.
PR 26-JUL-2000; 2000US-0220963.
PR 26-JUL-2000; 2000US-0220964.
PR 14-AUG-2000; 2000US-0224518.
PR 14-AUG-2000; 2000US-0224519.
PR 14-AUG-2000; 2000US-0225213.
PR 14-AUG-2000; 2000US-0225214.
PR 14-AUG-2000; 2000US-0225266.
PR 14-AUG-2000; 2000US-0225267.
PR 14-AUG-2000; 2000US-0225268.
PR 14-AUG-2000; 2000US-0225270.
PR 14-AUG-2000; 2000US-0225447.
PR 14-AUG-2000; 2000US-0225757.
PR 14-AUG-2000; 2000US-0225758.
PR 14-AUG-2000; 2000US-0225759.
PR 18-AUG-2000; 2000US-0226279.
PR 22-AUG-2000; 2000US-0226681.
PR 22-AUG-2000; 2000US-0226868.
PR 22-AUG-2000; 2000US-0227182.
PR 23-AUG-2000; 2000US-0227009.
PR 30-AUG-2000; 2000US-0228924.
PR 01-SEP-2000; 2000US-0229287.
PR 01-SEP-2000; 2000US-0229343.
PR 01-SEP-2000; 2000US-0229344.
PR 01-SEP-2000; 2000US-0229345.
PR 05-SEP-2000; 2000US-0229509.
PR 05-SEP-2000; 2000US-0229513.
PR 06-SEP-2000; 2000US-0230437.
PR 06-SEP-2000; 2000US-0230438.
PR 08-SEP-2000; 2000US-0231242.
PR 08-SEP-2000; 2000US-0231243.
PR 08-SEP-2000; 2000US-0231244.
PR 08-SEP-2000; 2000US-0231413.
PR 08-SEP-2000; 2000US-0231414.
PR 08-SEP-2000; 2000US-0232080.
PR 08-SEP-2000; 2000US-0232081.
PR 12-SEP-2000; 2000US-0231968.
PR 14-SEP-2000; 2000US-0232397.
PR 14-SEP-2000; 2000US-0232398.
PR 14-SEP-2000; 2000US-0232399.
PR 14-SEP-2000; 2000US-0232400.
PR 14-SEP-2000; 2000US-0232401.
PR 14-SEP-2000; 2000US-0232403.
PR 14-SEP-2000; 2000US-0233064.
PR 14-SEP-2000; 2000US-0233065.
PR 21-SEP-2000; 2000US-0234223.
PR 21-SEP-2000; 2000US-0234274.
PR 25-SEP-2000; 2000US-0234997.
PR 25-SEP-2000; 2000US-0234998.
PR 26-SEP-2000; 2000US-0235484.
PR 27-SEP-2000; 2000US-0235834.
PR 27-SEP-2000; 2000US-0235836.
PR 29-SEP-2000; 2000US-0236327.
PR 29-SEP-2000; 2000US-0236367.
PR 29-SEP-2000; 2000US-0236368.
PR 29-SEP-2000; 2000US-0236369.
PR 29-SEP-2000; 2000US-0236370.
PR 02-OCT-2000; 2000US-0236802.
PR 02-OCT-2000; 2000US-0237037.
PR 02-OCT-2000; 2000US-0237038.
PR 02-OCT-2000; 2000US-0237039.
PR 02-OCT-2000; 2000US-0237040.
PR 13-OCT-2000; 2000US-0239935.
PR 13-OCT-2000; 2000US-0239937.
PR 20-OCT-2000; 2000US-0240960.
PR 20-OCT-2000; 2000US-0241221.
PR 20-OCT-2000; 2000US-0241785.
PR 20-OCT-2000; 2000US-0241786.
PR 20-OCT-2000; 2000US-0241787.
PR 20-OCT-2000; 2000US-0241808.
PR 20-OCT-2000; 2000US-0241809.
PR 01-NOV-2000; 2000US-0241826.
PR 01-NOV-2000; 2000US-0244617.
PR 08-NOV-2000; 2000US-0246474.
PR 08-NOV-2000; 2000US-0246475.
PR 08-NOV-2000; 2000US-0246476.
PR 08-NOV-2000; 2000US-0246477.
PR 08-NOV-2000; 2000US-0246478.
PR 08-NOV-2000; 2000US-0246523.
PR 08-NOV-2000; 2000US-0246524.
PR 08-NOV-2000; 2000US-0246525.
PR 08-NOV-2000; 2000US-0246526.
PR 08-NOV-2000; 2000US-0246527.
PR 08-NOV-2000; 2000US-0246528.
PR 08-NOV-2000; 2000US-0246532.
PR 08-NOV-2000; 2000US-0246609.
PR 08-NOV-2000; 2000US-0246610.
PR 08-NOV-2000; 2000US-0246611.
PR 08-NOV-2000; 2000US-0246613.
PR 17-NOV-2000; 2000US-0249207.
PR 17-NOV-2000; 2000US-0249208.
PR 17-NOV-2000; 2000US-0249209.
PR 17-NOV-2000; 2000US-0249210.
PR 17-NOV-2000; 2000US-0249211.
PR 17-NOV-2000; 2000US-0249212.
PR 17-NOV-2000; 2000US-0249213.
PR 17-NOV-2000; 2000US-0249214.
PR 17-NOV-2000; 2000US-0249215.
PR 17-NOV-2000; 2000US-0249216.
PR 17-NOV-2000; 2000US-0249217.
PR 17-NOV-2000; 2000US-0249218.
PR 17-NOV-2000; 2000US-0249219.
PR 17-NOV-2000; 2000US-0249299.
PR 17-NOV-2000; 2000US-0249300.
PR 01-DEC-2000; 2000US-0250160.
PR 01-DEC-2000; 2000US-0250391.
PR 05-DEC-2000; 2000US-0251030.
PR 05-DEC-2000; 2000US-0251988.
PR 06-DEC-2000; 2000US-0256719.
PR 08-DEC-2000; 2000US-0251479.
PR 08-DEC-2000; 2000US-0251856.
PR 08-DEC-2000; 2000US-0251868.
PR 08-DEC-2000; 2000US-0251869.
PR 08-DEC-2000; 2000US-0251989.
PR 08-DEC-2000; 2000US-0251990.
PR 11-DEC-2000; 2000US-0254097.
PR 05-JAN-2001; 2001US-0259678.
PR (HUMA-) HUMAN GENOME SCI INC.
PA Rosen CA, Barash SC, Ruben SM;
PI WPI; 2001-488786/53.
XX P-PSDB; ABB10965.
XX

PT New isolated ovarian and/or breast cancer related nucleic acids and
PT polypeptides, useful for diagnosing, treating and/or preventing human
PT diseases and disorders, particularly ovarian and/or breast cancer -
XX
PS Claim 1: SEQ ID NO 233; 577pp + Sequence Listing; English.

The invention relates to novel genes (ABA07454-ABA08224) and proteins (AB01743-AB01980) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal and parasitic infections.

SQ Sequence 699 BP; 202 A; 161 C; 120 G; 211 T; 5 other;

[illegible]

RESULT 11	
AAI02390	
AAI02590 standard; cDNA; 699 bp.	
XX	
AC	AAI02590;
XX	
DT	21-NOV-2001 (first entry)
XX	
DE	Human reproductive system related antigen cDNA SEQ ID NO: 2591.
XX	
XX	Human; reproductive system related antigen; reproductive system disorder;
KW	cancer; gene therapy; ss.
XX	
OS	Homo sapiens.
XX	
PN	WO200155320-A2.
XX	
PD	02-AUG-2001.
XX	
PF	17-JAN-2001; 2001WO-US01339.
XX	
PR	31-JAN-2000; 2000US-0179065.
PR	04-FEB-2000; 2000US-0180628.
PR	24-FEB-2000; 2000US-0184664.
PR	02-MAR-2000; 2000US-0186350.
PR	16-MAR-2000; 2000US-0189874.
PR	17-MAR-2000; 2000US-0190076.
PR	18-APR-2000; 2000US-0198123.
PR	19-MAY-2000; 2000US-0205515.

CC chemokines, endogenously produced specific and non-specific enzymes,
CC binding proteins, adhesion molecules and their receptors, cytokine and
CC chemokine receptors, adenosine receptors, bradykinin receptors, central
CC nervous system (CNS) and peripheral nervous and non-nervous system
CC receptors, CNS and peripheral nervous and non-nervous system peptide
CC transmitters, defensins, growth factors, vasoactive peptides and
CC receptors, binding proteins and malignancy associated proteins. The
CC antisense oligonucleotides may be used in this way to treat disorders
CC including respiratory obstruction (especially pulmonary obstruction
CC and/or bronchoconstriction) and/or lung inflammation, allergies
CC and/or surfactant hypoproduction which are associated with a disease or
CC condition selected from pulmonary vasoconstriction, inflammation,
CC allergies, asthma, impaired respiration, respiratory distress syndrome
CC (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary
CC hypertension, emphysema, chronic obstructive pulmonary disease (COPD),
CC pulmonary transplantation rejection, pulmonary infections, bronchitis,
CC and/or cancer. AAF18434 to AAF21543 represent human polynucleotide
CC fragments and antisense oligonucleotides used in the exemplification of
CC the present invention.
XX
SQ Sequence 14968 BP; 5108 A; 2436 C; 2758 G; 4666 T; 0 other;

Query Match 7.2%; Score 72.4; DB 21; Length 14968;
Best Local Similarity 65.1%; Pred. No. 9.le-09;
Matches 125; Conservative 0; Mismatches 61; Indels 6; Gaps 1;
Qy 240 TTAATAAGGACCTTAGGGTGGTCTTAATCCCAATCTAAGTGATGTCTCCATGAAGAG 299
Db 6085 TTAATAAGGATACAAATGTGCACAGAGAGAAATGGCCACATGAGGACACAAATGAGAAT 359
Qy 300 GAAATAGGATACAAATGTGCACAGAGAGAAATGGCCACATGAGGACACAAATGAGAAT 359
Db 6145 GATAAGGCTATAGAT-----ACACAGAGATAGAAAACCATGTGAAGACACTGGGAGAAA 6198
Qy 360 GTGGCTACTTACAGCCTTAGGAGAGAGAGCCCTCCGAGAAACACACCCCTACCCACACTTG 419
Db 6199 ATGGCCACCTTATAGCCCAAGGAGGAGGACTCCAAAGAAACCAACACTGCTGGTACCCTG 6258
Qy 420 ATGTTGGACTTC 431
Db 6259 ATTCAGACCTC 6270

RESULT 15
ID AAA35221 standard; DNA; 14968 BP.
XX
AC AAA35221;
XX
XX 28-JUL-2000 (first entry)
DE Human adenosine receptor related polynucleotide 2nd SEQ ID NO:95.
XX
KW Human; adenosine receptor; low adenosine antisense oligonucleotide;
KW phosphorothioate; impaired respiration; inflammation; allergy;
KW allergic disease; bronchoconstriction; inhibitor; antiinflammatory;
KW antiallergic; antisthmatic; cytostatic; analgesic; impaired airway;
KW lung disease; ischaemic condition; pulmonary vasoconstriction; asthma;
KW respiratory distress syndrome; pain; cystic fibrosis; emphysema;
KW pulmonary hypertension; chronic obstructive pulmonary disease; COPD;
KW cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
XX
OS Homo sapiens.
XX
PN WO200009525-A2.
XX
PD 24-FEB-2000.
XX
PF 03-AUG-1999; 99WO-US17712.
XX
PR 03-AUG-1998; 98US-0095212.
XX
PA (UYEC-) UNIV EAST CAROLINA.

XX Nyce JW;
PI
XX WPI; 2000-205971/18.
DR
XX New antisense oligonucleotides useful for treating e.g. pulmonary
PT vasoconstriction, inflammation, allergies, asthma, hypertension,
PT bronchitis, emphysema, respiratory distress syndrome, ischemia or
PT cancers -
XX
PS Disclosure; Page 1260-1264; 1343pp; English.
XX
CC The present invention describes a new composition comprising an
CC antisense oligonucleotide (ON) with low adenosine (up to 15%), which
CC targets nucleic acids involved in bronchoconstriction, allergies, and/or
CC inflammation. The ON can have antiinflammatory, antiallergic,
CC antiasthmatic, cytostatic and analgesic activities. The compositions are
CC useful for the treatment of diseases associated with inflammation,
CC impaired airways, including lung disease and diseases whose secondary
CC effects afflict the lungs of a subject. They can be used for treating
CC e.g. ischaemic conditions, pulmonary vasoconstriction, allergies,
CC asthma, impaired respiration, respiratory distress syndrome, pain, cystic
CC fibrosis, pulmonary hypertension, emphysema, chronic obstructive
CC pulmonary disease (COPD), and cancers such as leukaemias, lymphomas,
CC carcinomas, and cancers which may metastasize to the lungs, including
CC breast and prostate cancer. The reduction of the adenosine content of
CC the ONs reduces side effects. The A-containing ONs break down with the
CC release of deoxyadenosine which activates adenosine receptors causing
CC bronchoconstriction and inflammation. AAA32313 to AAA35312 represent the
CC nucleotide sequences given in the sequence listing from the present
CC invention, which correspond to SEQ ID NO:1 to 2815, and then the last
CC 185 sequences are also called SEQ ID NO:1 to 185, but the sequences
CC differ from the previously named sequences. SEQ ID NO:11 to 1680
CC (AAA32323 to AAA33992) are specifically claimed ONs from the present
CC invention. N.B. Sequences given in the disclosure of the present
CC invention do not match up with their corresponding SEQ ID NO: sequences
CC given in the sequence listing.
XX
SQ Sequence 14968 BP; 5109 A; 2436 C; 2757 G; 4666 T; 0 other;

Query Match 7.2%; Score 72.4; DB 21; Length 14968;
Best Local Similarity 65.1%; Pred. No. 9.le-09;
Matches 125; Conservative 0; Mismatches 61; Indels 6; Gaps 1;
Qy 240 TTAATAAGGACCTTAGGGTGGTCTTAATCCCAATCTAAGTGATGTCTCCATGAAGAG 299
Db 6085 TTAATAAGGATACAAATGTGCACAGAGAGAAATGGCCACATGAGGACACAAATGAGAAT 359
Qy 300 GAAATAGGATACAAATGTGCACAGAGAGAAATGGCCACATGAGGACACAAATGAGAAT 359
Db 6145 GATAAGGCTATAGAT-----ACACAGAGATAGAAAACCATGTGAAGACACTGGGAGAAA 6198
Qy 360 GTGGCTACTTACAGCCTTAGGAGAGAGAGCCCTCCGAGAAACACACCCCTACCCACACTTG 419
Db 6199 ATGGCCACCTTATAGCCCAAGGAGGAGGACTCCAAAGAAACCAACACTGCTGGTACCCTG 6258
Qy 420 ATGTTGGACTTC 431
Db 6259 ATTCAGACCTC 6270

Search completed: December 24, 2002, 17:11:02
Job time : 841.952 secs